

# Symbiont ecology and its influence on the physiology and isotopic composition of a reef coral across space and time

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9/19/2018

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## Project background

### Sea level correction

```
#####  
# Sea level correction  
#####  
  
#### attach data files  
data<-read.csv("data/mastersheet_PanKBAY.csv") # master file  
qPCR.Innis<-read.csv("data/PanKBay_summer_qPCR.csv") # qPCR from summer (Innis et al. 2018)  
  
JuneTide=read.csv("data/environmental/sea level/Station_1612480_tide_ht_20160601-20160630.csv")  
JulyTide=read.csv("data/environmental/sea level/Station_1612480_tide_ht_20160701-20160731.csv")  
AugustTide=read.csv("data/environmental/sea level/Station_1612480_tide_ht_20160801-20160831.csv")  
DecemberTide=read.csv("data/environmental/sea level/Station_1612480_tide_ht_20161201-20161222.csv")  
  
Tide<-rbind(JuneTide, JulyTide, AugustTide, DecemberTide) # all MSL in meters  
  
Tide$Time <- as.POSIXct(Tide$TimeUTC, format="%Y-%m-%d %H:%M:%S", tz="UTC")  
attributes(Tide$Time)$tzone <- "Pacific/Honolulu"  
# use attr(as.POSIXlt(Tide$Time),"tzone") to confirm in PST  
  
Tide<-Tide[, c(-5:-8)] #remove unnecessary columns
```

```

data$date.time<- as.POSIXct(paste(as.character(data$Date), as.character(data$Time.of.collection)),
                             format="%m/%d/%y %H:%M", tz="Pacific/Honolulu") # make sure time in HST for master

data$Time.r<-round_date(data$date.time,unit="6 minutes")
Tide$Time.r <- Tide$Time
data<-merge(data, Tide, by="Time.r", all.x=T)
data$newDepth <- data$Depth..m-data$TideHT # new depth in METERS

```

## Light at Depth

```

##### # LIGHT AT DEPTH
#####
#####
# using new depth and the site-specific kd (light attenuation) determine approximate "light at depth" f

#####

### attach necessary files
logger.depths<-read.csv("data/environmental/temp and light/light.logger.depths.csv") # depths for logger
kd.all<-read.csv("data/environmental/kd.all.csv") # kds for each site

# Seasonal DLI used for "period of collection" light levels
month.DLI<-read.csv("data/environmental/temp and light/Jun_DecPAR/All.DLI_long.csv")

# corals collected in June-July-August use summer time DLI for these months as indicator of average DLI
summer.DLI<-month.DLI[(month.DLI$Month=="June" | month.DLI$Month=="July" | month.DLI$Month=="August"),]
winter.DLI<-month.DLI[(month.DLI$Month=="November" | month.DLI$Month=="December" | month.DLI$Month=="January"),]

# summer mean and SE dataframe
sum.mean<-aggregate(DLI~Site, summer.DLI, mean)
sum.SE<-aggregate(DLI~Site, summer.DLI, std.error)
sum.light.df<-cbind(sum.mean, sum.SE[2])
colnames(sum.light.df)=c("Site", "mean.DLI", "SE")
sum.light.df$Season<-as.factor("summer")

# winter mean and SE dataframe
wint.mean<-aggregate(DLI~Site, winter.DLI, mean)
wint.SE<-aggregate(DLI~Site, winter.DLI, std.error)
wint.light.df<-cbind(wint.mean, wint.SE[2])
colnames(wint.light.df)=c("Site", "mean.DLI", "SE")
wint.light.df$Season<-as.factor("winter")

season.DLI<-rbind(sum.light.df[,c(4,1,2:3)], wint.light.df[,c(4,1,2:3)]) # compiled means for DLI at ~2m

write.csv(season.DLI, "data/environmental/season.DLI.csv")
#####
### make new dataframe for calculations
df.light<- data[, c("Season", "Location", "newDepth")]

# make a column of "depth differences" relative to where ~2m logger was deployed
df.light$depth.diff<-ifelse(df.light$Location=="F1-R46", df.light$newDepth-1.8288,
                           ifelse(df.light$Location=="F8-R10", df.light$newDepth-1.8288,

```

```

        ifelse(df.light$Location=="HIMB", df.light$newDepth-1.8288,
              df.light$newDepth-2.4384))) # last statement for remaining site

# make a column for sample-specific light at depth (estimate) based on kd
# follow: #with 2m as baseline#  $E(\text{depth}) = E(2m) \cdot \exp(-k_d \cdot (\text{depth} - 2m))$ 
# so that: light at depth = DLI at mid.depth * exp (-kd *(delta shallow-deep in m))

df.light$Light<-ifelse(df.light$Location=="HIMB" & df.light$Season=="summer",
                      season.DLI$mean.DLI[1]*exp(-kd.all$kd[1]*df.light$depth.diff), # summer HIMB
                      ifelse(df.light$Location=="F8-R10" & df.light$Season=="summer",
                              season.DLI$mean.DLI[2]*exp(-kd.all$kd[2]*df.light$depth.diff), # summer R10
                              ifelse(df.light$Location=="R42" & df.light$Season=="summer",
                                      season.DLI$mean.DLI[3]*exp(-kd.all$kd[3]*df.light$depth.diff), # summer R42
                                      ifelse(df.light$Location=="F1-R46" & df.light$Season=="summer",
                                              season.DLI$mean.DLI[4]*exp(-kd.all$kd[4]*df.light$depth.diff), # summer R46
                                              ))))
                      ifelse(df.light$Location=="HIMB" & df.light$Season=="winter",
                              season.DLI$mean.DLI[5]*exp(-kd.all$kd[1]*df.light$depth.diff), # winter HIMB
                              ifelse(df.light$Location=="F8-R10" & df.light$Season=="winter",
                                      season.DLI$mean.DLI[6]*exp(-kd.all$kd[2]*df.light$depth.diff), # winter R10
                                      ifelse(df.light$Location=="R42" & df.light$Season=="winter",
                                              season.DLI$mean.DLI[7]*exp(-kd.all$kd[3]*df.light$depth.diff), # winter R42
                                              season.DLI$mean.DLI[8]*exp(-kd.all$kd[4]*df.light$depth.diff)) # winter R46
                              ))))

##### plot of light x depth by season
df.light$Location <- factor(df.light$Location, levels = c("F1-R46", "R42", "F8-R10", "HIMB"))

plot.by.sites=ggplot(df.light, aes(Light)) + geom_density(aes(fill=Location), alpha=0.3, position = 'stack')

plot.by.site=ggplot(df.light, aes(Light)) + geom_density(aes(fill=Season), alpha=0.3, position = 'stack')
  scale_x_continuous(limits=c(0, 40)) + ggtitle("Light at Depth by Seasons") + facet_wrap(~Location, scales="free")

#####
# can loop as a list
p=ggplot(df.light, aes(Light, fill=Season)) + geom_density(alpha=0.3, position = 'stack') + scale_x_continuous(limits=c(0, 40))

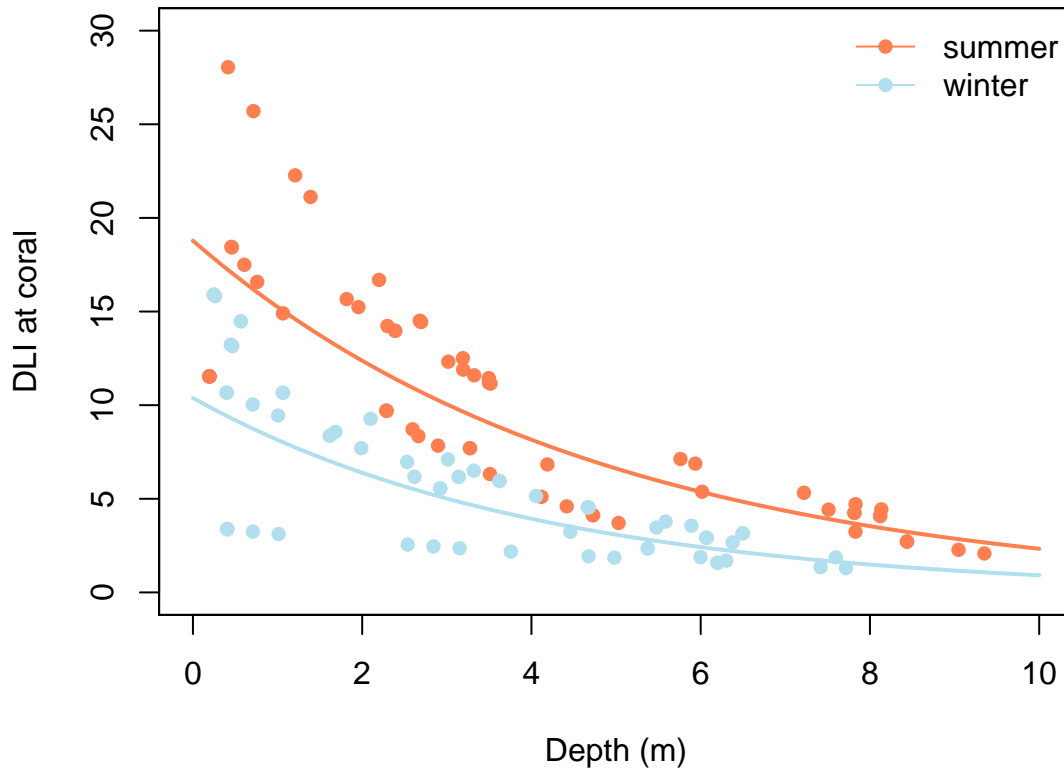
plots=dplyr::dfply(df.light, .(Location), function(x) p %>% x + facet_wrap(~Location))

##### plot of light x depth by season with exponential curve fitting
Sum<-df.light[(df.light$Season=="summer"),]
Win<-df.light[(df.light$Season=="winter"),]

plot(Light~newDepth, Sum, col="coral", pch=16, xlab="Depth (m)", ylab="DLI at coral",
     ylim=c(0, 30), xlim=c(0, 10))
summod<-lm(log(Light)~newDepth, Sum)
curve(exp(coef(summod)["(Intercept)"]+coef(summod)["newDepth"]*x), add=TRUE, col="coral", lwd=2)
par(new=T)
plot(Light~newDepth, Win, col="lightblue2", pch=16, xaxt="n", yaxt="n",
     xlab="", ylab="", ylim=c(0, 30), xlim=c(0, 10))
wintmod<-lm(log(Light)~newDepth, Win)

```

```
curve(exp(coef(wintmod)["(Intercept)"]+coef(wintmod)["newDepth"]*x), add=TRUE, col="lightblue2", lwd=2)
legend("topright", legend=c("summer", "winter"), col=c("coral", "lightblue2"), lty=1, lwd=1, pch=16, bt
```



### Isotopes in particulates

Plankton tows and seawater collections to parameterized particulates and plankters as potential heterotrophic end members available to reef corals. Sampling was done at 4 sites where corals were collected [*North*: (Reef 42, fringe-Reef 46), and *South*: (HIMB, fringe-Reef 10)], as well as two reefs in central region of the bay where corals were not collected (*Central*: Reef 25, fringe-Reef 25) to increase spatial resolution of suspended particulate isotope sample sizes. Using size-fractionated materials collected in seawater and plankton tows, we examined the spatial (*among reef sites*) and temporal patterns (*among seasons*) in stable isotope values of size-fractionated end members. Carbon and nitrogen isotope values among the 6 reef sites were not significantly different ( $p > 0.140$ ), season had marginal effects ( $p > 0.049$ ), whereas fraction influenced isotope values significantly ( $p < 0.001$ ). Therefore, isotope values were pooled among reefs and seasons to best interpret size-fraction isotope values.

```
SWiso<-read.csv("data/isotopes_SW_all times.csv")
```

```
anova(lm(d13C~Reef.ID+Time.point+SW.fraction..um, data=SWiso)) # no site effect
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: d13C
```

```
##      Df  Sum Sq Mean Sq F value    Pr(>F)
## Reef.ID      5   17.926   3.5852    1.3418 0.2626013
## Time.point    1    7.921    7.9207    2.9645 0.0914188 .
## SW.fraction..um 4   76.419   19.1048    7.1504 0.0001286 ***
## Residuals    49  130.920    2.6718
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

anova(lm(d15N~Reef.ID+Time.point+SW.fraction..um, data=SWiso)) # no site effect

## Analysis of Variance Table
##
## Response: d15N
##          Df Sum Sq Mean Sq F value    Pr(>F)
## Reef.ID    5  2.3255   0.4651   1.7291   0.14557
## Time.point  1  1.0935   1.0935   4.0653   0.04927 *
## SW.fraction..um 4 30.3773   7.5943  28.2335  3.454e-12 ***
## Residuals   49 13.1802   0.2690
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

SWiso$Reef.ID<-factor(SWiso$Reef.ID, levels=c("F1-46", "R42", "F2-R25", "R25", "F8-R10", "HIMB"))
SWiso$SW.fraction..um<-factor(SWiso$SW.fraction..um, levels=c(">243", "<243", "100-243", "10-100", "0-100"))

winter.data<-SWiso[(SWiso$Time.point=="winter"),]
summer.data<-SWiso[(SWiso$Time.point=="summer"),]

# plots of SW isotopes by Season-- first showing by size, then by site
op<-par(mfrow = c(2,2), mar=c(5,5,2,1))

##### Sizes
# Summer size d13C
plot(d13C~SW.fraction..um, data=summer.data, ylim=c(-25,-10),
     main=expression(paste("summer"~ delta^{13}, "C")), ylab=expression(paste(delta^{13}, "C (%, V-PDB)")),
     col="paleturquoise3", cex.axis=0.7, cex.main=1, cex.lab= 0.8, xlab=expression(paste("Size Fractionation (%, V-PDB)")))

# Winter size d13C
plot(d13C~SW.fraction..um, data=winter.data, ylim=c(-25,-10),
     main=expression(paste("winter"~ delta^{13}, "C")), ylab=expression(paste(delta^{13}, "C (%, V-PDB)")),
     col="paleturquoise3", cex.axis=0.7, cex.main=1, cex.lab= 0.8, xlab=expression(paste("Size Fractionation (%, V-PDB)")))

# Summer size d15N
plot(d15N~SW.fraction..um, data=summer.data, ylim=c(3,10),
     main=expression(paste("summer"~ delta^{15}, "N")), col="plum", cex.axis=0.7, cex.main=1, cex.lab= 0.8,
     xlab=expression(paste("Size Fractionation (%, air)")),
     ylab=expression(paste(delta^{15}, "N (%, air)")))

# Winter size d15N
plot(d15N~SW.fraction..um, data=winter.data, ylim=c(3,10),
     main=expression(paste("winter"~ delta^{15}, "N")), ylab=expression(paste(delta^{15}, "N (%, air)")),
     col="plum", cex.axis=0.7, cex.main=1, cex.lab= 0.8, xlab=expression(paste("Size Fractionation (%, air)")))

##### Sites
op<-par(mfrow = c(2,2), mar=c(5,5,2,1))

# Summer site d13C
plot(d13C~Reef.ID, data=summer.data, ylim=c(-25,-10),
     main=expression(paste("summer"~ delta^{13}, "C")), ylab=expression(paste(delta^{13}, "C (%, V-PDB)")),
     col="lightskyblue", cex.axis=0.7, cex.main=1, cex.lab= 0.8, xlab="Reef Sites")

# Winter site d13C
plot(d13C~Reef.ID, data=winter.data, ylim=c(-25,-10),
```

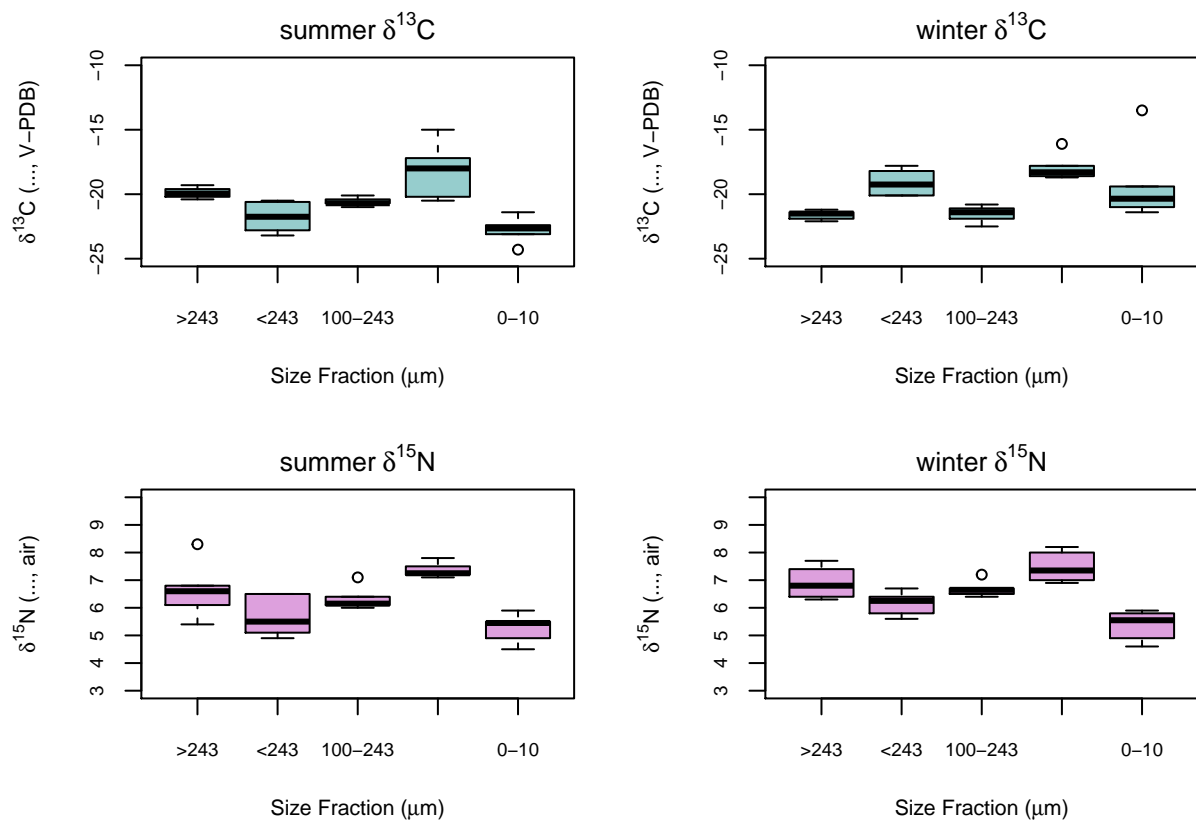


Figure 1: *Figures of size-fractionated sample isotope values stratified by seasons (summer and winter)*

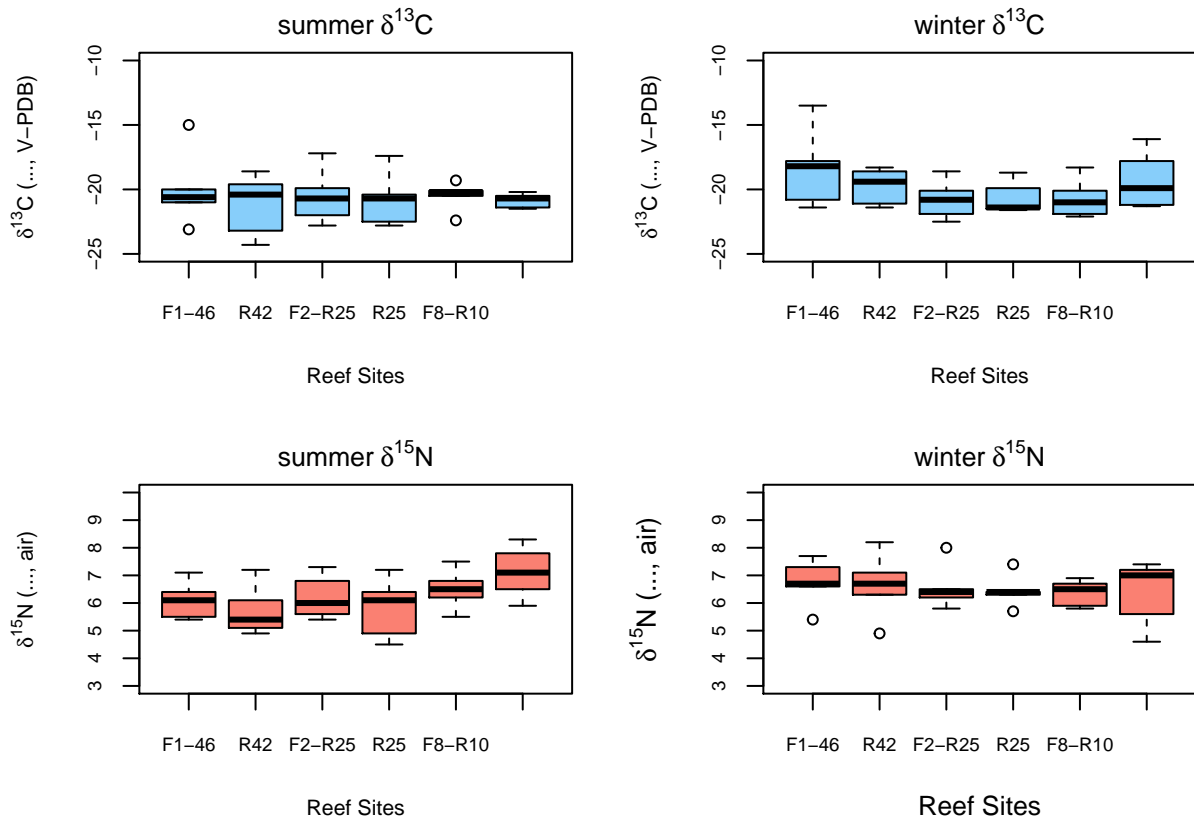


Figure 2: *Figures of isotope values in seawater particles stratified by sites (northwest to southeast)*

```

main=expression(paste("winter"~ delta^{13}, "C")), ylab=expression(paste(delta^{13}, "C (‰, V-PDB)
col="lightskyblue", cex.axis=0.7, cex.main=1, cex.lab= 0.8, xlab="Reef Sites")

# Summer site d15N
plot(d15N~Reef.ID, data=summer.data, ylim=c(3,10),
     main=expression(paste("summer"~ delta^{15}, "N")), col="salmon", cex.axis=0.7, cex.main=1, cex.lab=
     ylab=expression(paste(delta^{15}, "N (‰, air)")), xlab="Reef Sites")

# Winter site d15N
plot(d15N~Reef.ID, data=winter.data, ylim=c(3,10),
     main=expression(paste("winter"~ delta^{15}, "N")), ylab=expression(paste(delta^{15}, "N (‰, air)")),
     col="salmon", cex.axis=0.7, cex.main=1, xlab="Reef Sites")

####
#### making scatter for d15N and d13C, pooled across seasons and sites
mix.N.mean<-aggregate(d15N~SW.fraction..um, data=SWiso, mean)
mix.N.SE<-aggregate(d15N~SW.fraction..um, data=SWiso, std.error)
mix.C.mean<-aggregate(d13C~SW.fraction..um, data=SWiso, mean)
mix.C.SE<-aggregate(d13C~SW.fraction..um, data=SWiso, std.error)
mix.data<-cbind(mix.N.mean, mix.C.mean[c(2,0)], mix.N.SE[c(2,0)], mix.C.SE[c(2,0)]); colnames(mix.data)=

colors=c("#FF6A6A", "#00B2EE", "#FFB90F", "#3CB371", "#8B7500")
op<-par(mfrow = c(1,1), mar=c(5,4,1,5),xpd=TRUE, pty="sq")

```

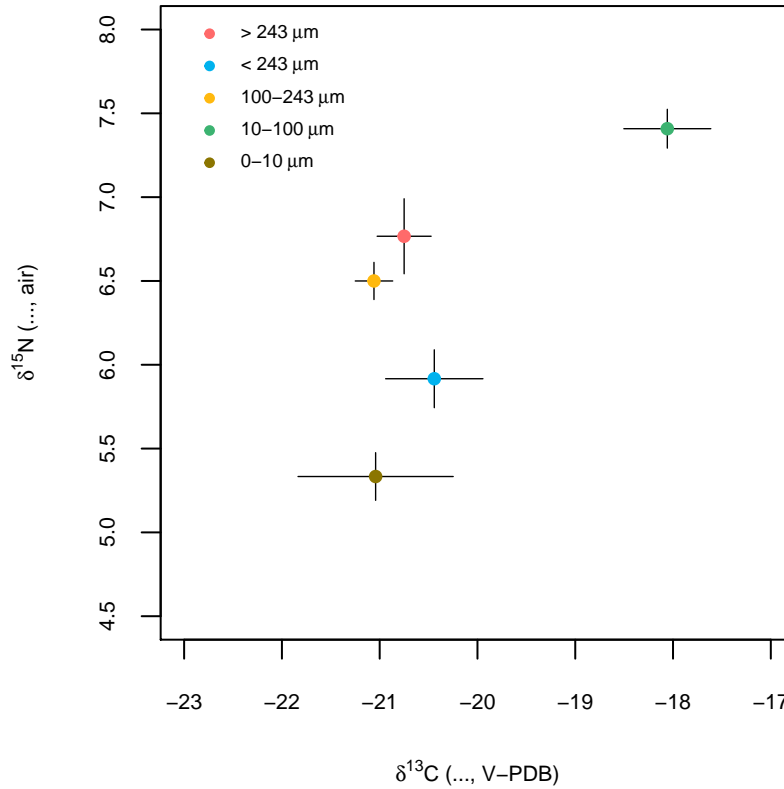


Figure 3: Isotope values for size-fractionated seawater particles and plankters pooled across 2 seasons and 6 reef sites

```
size.labels=c(expression(paste("> 243"~mu,m)), expression(paste("< 243"~mu,m)), expression(paste("100-243"~mu,m)),
expression(paste("10-100"~mu,m)), expression(paste("0-10"~mu,m)))

#### make the plot
plot(d15N~d13C, data=mix.data, type="n", ylim=c(4.5,8), xlim=c(-23,-17), tck=-0.03, cex.axis=0.7, cex.lab=1.2,
      ylab=expression(paste(delta^{15}, "N (‰, air)")), xlab=expression(paste(delta^{13}, "C (‰, V-PDB)")),
      legend("topleft", inset=c(0.05,0.0), legend=size.labels, col=colors, pch=19, cex=0.6, bty="n", x.intersp=1.5),
      arrows(mix.data$d13C-mix.data$d13C.SE, mix.data$d15N, mix.data$d13C+mix.data$d13C.SE, mix.data$d15N, length=0.05, col="black"),
      arrows(mix.data$d13C, mix.data$d15N-mix.data$d15N.SE, mix.data$d13C, mix.data$d15N+mix.data$d15N.SE, length=0.05, col="black"),
      points(d15N~d13C, data=mix.data, pch=19, cex=0.8, col=colors))

dev.copy(pdf, "output/iso.sources.KBay.pdf", encod="MacRoman", height=4, width=4)
dev.off()
```

## Biology

```
#####
### Biological responses
#####

#data : this is the master file

# add in light at depth column from df.light dataframe
data$Light<-df.light$Light
```



```
##### produce a categorical depth bin #####
depth<-data$newDepth
data$depth.bin<-factor(ifelse(depth<2, "<2m", ifelse(depth >2 & depth <4, "2-4m", ifelse(depth >4 & dep

aggregate(Sample.ID~depth.bin+Season+Location, data, length)
data$depth.bin.small<-factor(ifelse(depth<4, "<4m", ">4m"), levels= c("<4m", ">4m"))

#####
# calculate, normalized dependent variables
#####
str(data)
data$cells.ml<-as.numeric(data$cells.ml)

# helpful shorthand
SA<-data$surface.area # surface area in cm2
blastate<-data$total.blastate.ml # tissue slurry blastate in ml

# AFDW.mg. == convert AFDW g to mg, mutiply by blastate volume, divide by cm2
data$biomass<- (data$mg.biomass.ml*blastate)/SA

# Symbiodinium.cells. == cell.ml * blastate / SA
data$zoox<- (data$cells.ml*blastate)/SA

# total chlorophyll == ug.chl.a.ml * blastate + ug.chl.c2.ml * blastate / SA
data$chltot<-(data$ug.chl.a.ml)+(data$ug.chl.c2.ml)*blastate/SA

# pg.chlorophyll.a..cell + pg.chlorophyll.c2..cell == ug.chltot.ml * 10^6 / cells.ml
data$chlcell<- (data$ug.chl.a.ml*10^6+data$ug.chl.c2.ml*10^6)/data$cells.ml
```

## qPCR

```
#####
# qPCR
#####

# qPCR
# Use steponeR to import data and calculate proporation of C and D symbionts
source_url("https://raw.githubusercontent.com/jrcunning/steponeR/master/steponeR.R")
Mcap.plates <- list.files(path="data/qPCR", pattern = "txt$", full.names = T); Mcap.plates
Mcap <- steponeR(files=Mcap.plates, delim="\t",
                 target.ratios=c("C.D"),
                 fluor.norm=list(C=2.26827, D=0),
                 copy.number=list(C=33, D=3),
                 ploidy=list(C=1, D=1),
                 extract=list(C=0.813, D=0.813))

Mcap <- Mcap$result
head(Mcap)

# remove +/-control
Mcap <- Mcap[grep("+C52", Mcap$Sample.Name, fixed=T, invert = T), ]
Mcap <- Mcap[grep("H20", Mcap$Sample.Name, fixed=T, invert = T), ]
```

```

# to remove any early-amplification CT noise
Mcap$C.CT.mean[which(Mcap$C.CT.mean < 15)] <- 0

#Remove failed samples, i.e., those where either C or D were NOT found in both reps
Mcap$fail <- ifelse(Mcap$C.reps < 2 & Mcap$D.reps < 2, TRUE, FALSE)
fails <- Mcap[Mcap$fail==TRUE, ]
Mcap <- Mcap[which(Mcap$fail==FALSE),]

# replace CT means with 'NA' as zero
Mcap$C.CT.mean[is.na(Mcap$C.CT.mean)] <-0
Mcap$D.CT.mean[is.na(Mcap$D.CT.mean)] <-0

Mcap$C.D[is.na(Mcap$C.D)] <- 1 # sets all infinity (= 100% C) to 1.0

# calculate proportion C and proportion D where C and D are both present
Mcap$propC<- Mcap$C.D / (Mcap$C.D + 1)
Mcap$propD<- 1 / (Mcap$C.D + 1)

# where C and D are not cooccurring...
# if C.D = 1 = 100% C, make 'PropC' = 1 and 'PropD' = 0
# if C.D = 0 = 100% D, make 'PropD' = 1 and 'PropC' = 0
Mcap$propC[which(Mcap$C.D==1)] <- 1
Mcap$propD[which(Mcap$propC==1)] <- 0
Mcap$propD[which(Mcap$C.D==0)] <- 1

# calculate FOUR COMMUNITY categories: C, C>D, D>C, D
Mcap$Mix <- factor(ifelse(Mcap$propC > Mcap$propD, ifelse(Mcap$propD!= 0, "CD", "C"), ifelse(Mcap$propD
# Identify SINGLE dominant symbiont clade: C or D
Mcap$Dom <- factor(substr(as.character(Mcap$Mix), 1, 1))

# Set zeros to NA to facilitate log transformation
Mcap$propC[which(Mcap$propC==0)] <- NA
Mcap$propD[which(Mcap$propD==0)] <- NA

##### look for duplicates in dataset by year and type of event (bleach/recover)
Mcap[duplicated(Mcap$Sample.Name), ] ## duplicates

# remove duplicated
Mcap<-Mcap[!(Mcap$Sample.Name=="HIMB_15" & Mcap$File.Name=="Wall_PanKbay_plate1.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R10_05" & Mcap$File.Name=="Wall_PanKbay_plate1.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R42_06" & Mcap$File.Name=="Wall_PanKbay_plate1.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R46_01" & Mcap$File.Name=="Wall_PanKbay_plate1.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R46_02" & Mcap$File.Name=="Wall_PanKbay_plate2.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R46_03" & Mcap$File.Name=="Wall_PanKbay_plate1.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="HIMB_13" & Mcap$File.Name=="Wall_PanKbay_plate2.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="HIMB_14" & Mcap$File.Name=="Wall_PanKbay_plate2.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R10_15" & Mcap$File.Name=="Wall_PanKbay_plate3.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R10_15" & Mcap$File.Name=="Wall_PanKbay_plate1.txt"),]
Mcap<-Mcap[!(Mcap$Sample.Name=="R42_11" & Mcap$File.Name=="Wall_PanKbay_plate2.txt"),]

# parse Sample ID and Site from "Site.Name"
Mcap<-cbind(Mcap, colsplit(Mcap$Sample.Name, pattern= "_", c("Location", "Sample.ID")))

```

```

Mcap$Season<-as.factor("winter")
Mcap$Location<-as.factor(Mcap$Location)

Mcap$Location<-revalue(Mcap$Location, c("R10"="F8-R10", "R46"="F1-R46")) # rename factor levels

# make new factors for bay region and reef type
Mcap$Bay.region <- ifelse(Mcap$Location=="R42" | Mcap$Location=="F1-R46", "northern", "southern")
Mcap$Reef.type <- ifelse(Mcap$Location=="R42" | Mcap$Location=="HIMB", "patch", "fringe")

### reorder columns and finish
Mcap<-Mcap[, c(17,15,19,18,16, 1:14)] # reordered to match masterdata, and finish

### structure winter and summer qPCR dataframes to have same columns and combine dataframe
qPCR.winter<-Mcap[, (names(Mcap) %in%
  c("Season", "Location", "Reef.type", "Bay.region", "Sample.ID", "propC", "propD", "Mix", "D
qPCR.summer<-qPCR.Innis[, (names(qPCR.Innis) %in%
  c("Season", "Location", "Reef.type", "Bay.region", "Sample.ID", "propC", "propD", "Mix", "D

# merge qPCR files
qPCR.all<-rbind(qPCR.winter, qPCR.summer)

# add to master data
data.all<-merge(data, qPCR.all, by=c("Season", "Location", "Reef.type", "Bay.region", "Sample.ID"), all

##### remove columns no longer needed, update "Depth" to be tide-corrected depth )= newDepth
data.trim<-data.all[, !(names(data.all) %in% c("total.blastate.ml", "Date", "Time.of.collection", "Dep

data.trim$symb..C.N[data.trim$symb..C.N>=12.520270]=NA # set this outlier to NA

```

## Models

### physiology

#### Total Biomass

```

##### 
##### biomass ----
Y<-model.data$biomass
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
summary(full)

```

```

## Linear mixed model fit by REML t-tests use Satterthwaite approximations
##   to degrees of freedom [lmerMod]
## Formula: Y ~ Season * Light * Dom + (1 | Location)
##   Data: model.data
##
## REML criterion at convergence: 805.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.4430 -0.6942 -0.0371  0.5787  2.9820
##
## Random effects:
##   Groups   Name                Variance Std.Dev.

```

```
## Location (Intercept) 8.222 2.867
## Residual 56.407 7.510
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 28.7724 2.6187 19.3400 10.987 9.27e-10 ***
## Seasonwinter -7.6360 2.9066 108.3200 -2.627 0.00986 **
## Light -0.3483 0.2189 109.3900 -1.592 0.11437
## DomD -11.6171 6.6078 109.7400 -1.758 0.08152 .
## Seasonwinter:Light 1.0487 0.3636 109.6400 2.884 0.00473 **
## Seasonwinter:DomD 17.2336 7.9247 109.3400 2.175 0.03181 *
## Light:DomD 0.9354 0.4399 109.2600 2.126 0.03574 *
## Seasonwinter:Light:DomD -1.2701 0.6885 109.2600 -1.845 0.06779 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) Ssnwnt Light DomD Ssnw:L Ssn:DD Lgh:DD
## Seasonwintr -0.613
## Light -0.719 0.622
## DomD -0.254 0.258 0.252
## Ssnwntr:Lgh 0.406 -0.803 -0.565 -0.198
## Ssnwntr:DmD 0.209 -0.373 -0.206 -0.832 0.311
## Light:DomD 0.335 -0.324 -0.466 -0.919 0.318 0.763
## Ssnwnt:L:DD -0.208 0.425 0.290 0.594 -0.533 -0.857 -0.643
```

```
print(anova(full, type=2), digits=4)
```

```
## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season 215.45 215.45 1 108.2 3.820 0.0532 .
## Light 53.34 53.34 1 109.8 0.946 0.3330
## Dom 0.30 0.30 1 109.4 0.005 0.9421
## Season:Light 289.47 289.47 1 109.2 5.132 0.0255 *
## Season:Dom 266.76 266.76 1 109.3 4.729 0.0318 *
## Light:Dom 86.12 86.12 1 108.8 1.527 0.2192
## Season:Light:Dom 191.95 191.95 1 109.3 3.403 0.0678 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
ranef(full)
```

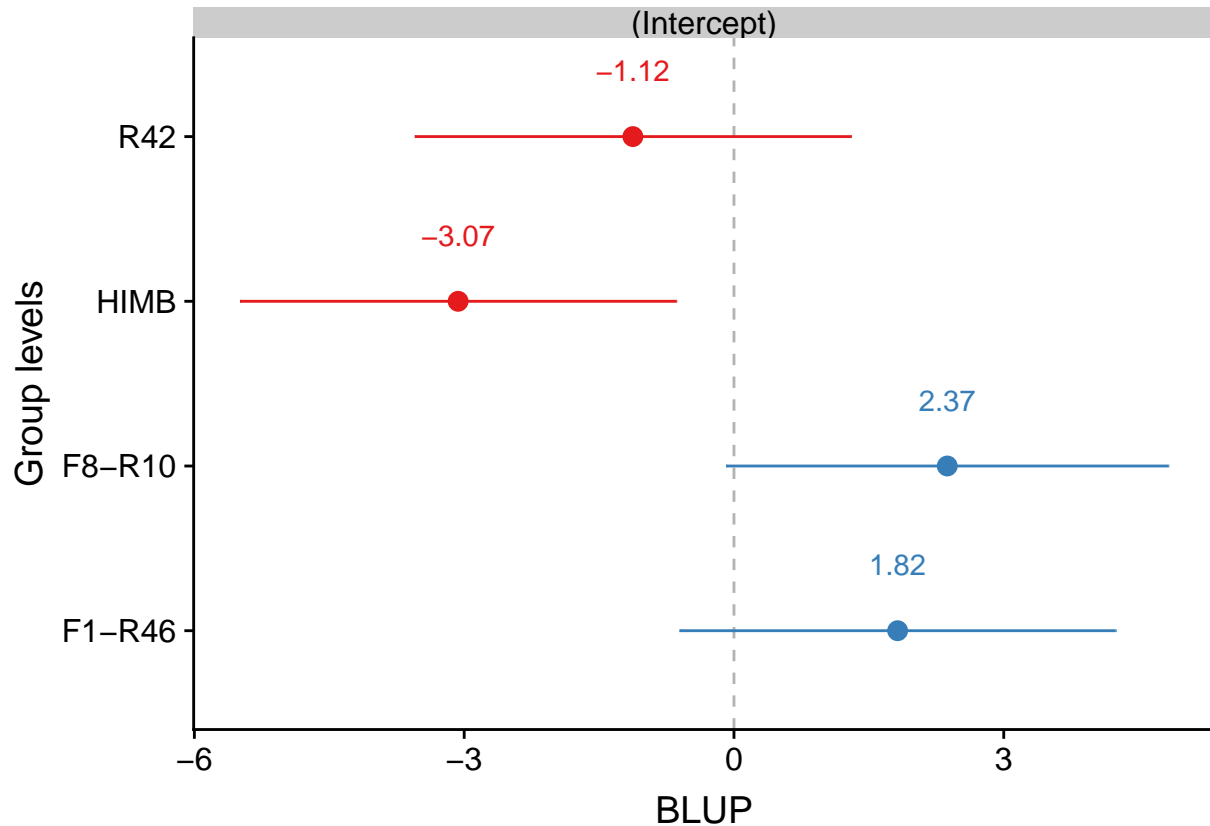
```
## $Location
## (Intercept)
## F1-R46 1.819827
## F8-R10 2.371571
## HIMB -3.067216
## R42 -1.124182
```

```
fixef(full)
```

```
## (Intercept) Seasonwinter Light
## 28.7724070 -7.6359780 -0.3483208
## DomD Seasonwinter:Light Seasonwinter:DomD
```

```
##          -11.6171087          1.0487075          17.2336137
##          Light:DomD Seasonwinter:Light:DomD
##          0.9353690          -1.2700733
```

```
sjp.lmer(full, y.offset = .4)
```



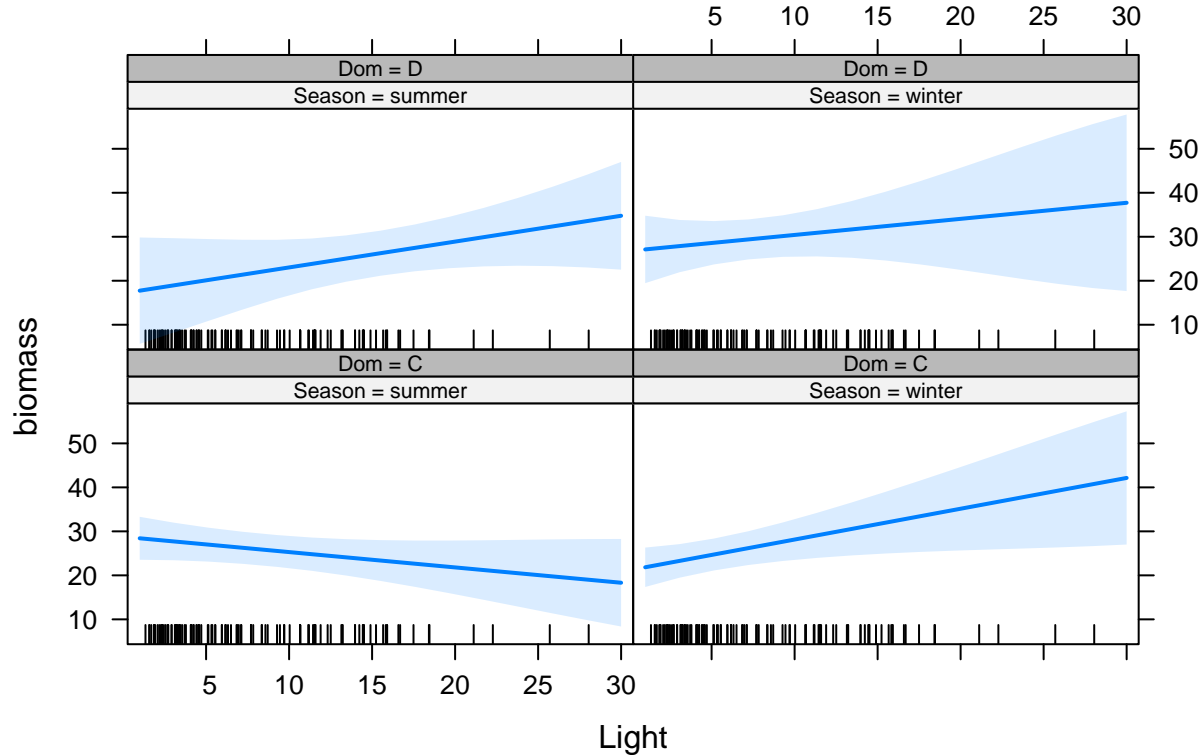
```
#sjp.lmer(full, vars = "Seasonwinter", type = "ri.slope")
```

```
posthoc<-emmeans(full, ~Light:Season)
CLD(posthoc, Letters=letters)
```

```
##      Light Season  emmean      SE    df lower.CL upper.CL .group
## 8.238607 summer 23.94724 2.302325 11.73 18.91804 28.97644   a
## 8.238607 winter 28.33614 1.844474  5.11 23.62535 33.04693   b
##
## Results are averaged over the levels of: Dom
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
## significance level used: alpha = 0.05
```

```
plot(allEffects(full), ylab="biomass", par.strip.text=list(cex=0.7))
```

## Season\*Light\*Dom effect plot



## Chlorophyll a

```
##### chltotal--
Y<-model.data$chltot
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
summary(full)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season * Light * Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 417.9
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.26787 -0.64400 -0.09381  0.53088  3.12573
##
## Random effects:
## Groups   Name                Variance Std.Dev.
## Location (Intercept) 0.3246   0.5697
## Residual              1.7029   1.3050
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    5.42443    0.47567 14.97000  11.404 8.82e-09
## Seasonwinter    2.64741    0.50510 108.18000   5.241 7.95e-07
```

```

## Light          -0.04057    0.03805 109.10000  -1.066  0.28869
## DomD           1.02243    1.14922 109.42000   0.890  0.37559
## Seasonwinter:Light -0.14846    0.06324 109.33000  -2.348  0.02070
## Seasonwinter:DomD -3.81872    1.37790 109.06000  -2.771  0.00656
## Light:DomD     -0.12052    0.07649 108.99000  -1.576  0.11799
## Seasonwinter:Light:DomD 0.27682    0.11971 108.98000   2.313  0.02263
##
## (Intercept)          ***
## Seasonwinter          ***
## Light
## DomD
## Seasonwinter:Light    *
## Seasonwinter:DomD     **
## Light:DomD
## Seasonwinter:Light:DomD *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) Ssnwnt Light  DomD   Ssnw:L Ssn:DD Lgh:DD
## Seasonwintr -0.586
## Light       -0.688  0.621
## DomD        -0.242  0.258  0.250
## Ssnwntr:Lgh  0.388 -0.803 -0.563 -0.200
## Ssnwntr:DmD  0.199 -0.374 -0.205 -0.832  0.312
## Light:DomD   0.320 -0.325 -0.464 -0.919  0.319  0.763
## Ssnwnt:L:DD -0.198  0.425  0.288  0.594 -0.533 -0.857 -0.644

print(anova(full, type=2), digits=4)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##          Sum Sq Mean Sq NumDF DenDF F.value  Pr(>F)
## Season      34.97   34.97     1 108.0  20.533 1.52e-05 ***
## Light       20.83   20.83     1 110.7  12.230 0.000678 ***
## Dom         10.85   10.85     1 109.1   6.370 0.013043 *
## Season:Light  3.07    3.07     1 108.9   1.801 0.182342
## Season:Dom    13.08   13.08     1 109.1   7.681 0.006564 **
## Light:Dom      0.03    0.03     1 108.6   0.016 0.900351
## Season:Light:Dom 9.11    9.11     1 109.0   5.348 0.022628 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

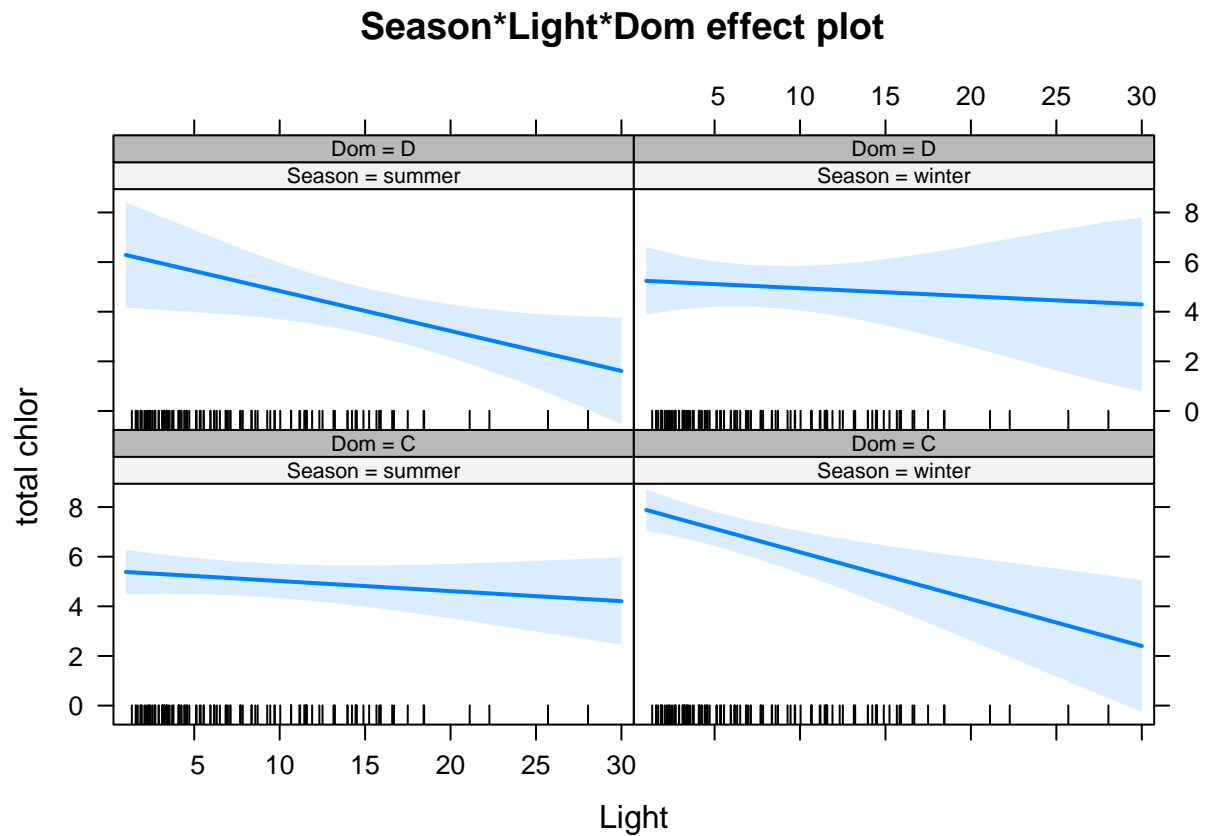
posthoc<-emmeans(full, ~Dom:Light:Season)
CLD(posthoc, Letters=letters)

## Dom   Light Season  emmean      SE    df lower.CL upper.CL .group
## D    8.238607 winter 5.005924 0.4237749  9.88 4.060098 5.951751  a
## C    8.238607 summer 5.090166 0.3452653  4.51 4.172822 6.007509  a
## D    8.238607 summer 5.119654 0.6589125 42.24 3.790143 6.449166  ab
## C    8.238607 winter 6.514512 0.3843448  6.70 5.597381 7.431643  b
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates

```

```
## significance level used: alpha = 0.05
```

```
plot(allEffects(full), ylab="total chlor", par.strip.text=list(cex=0.7))
```



```
ranef(full)
```

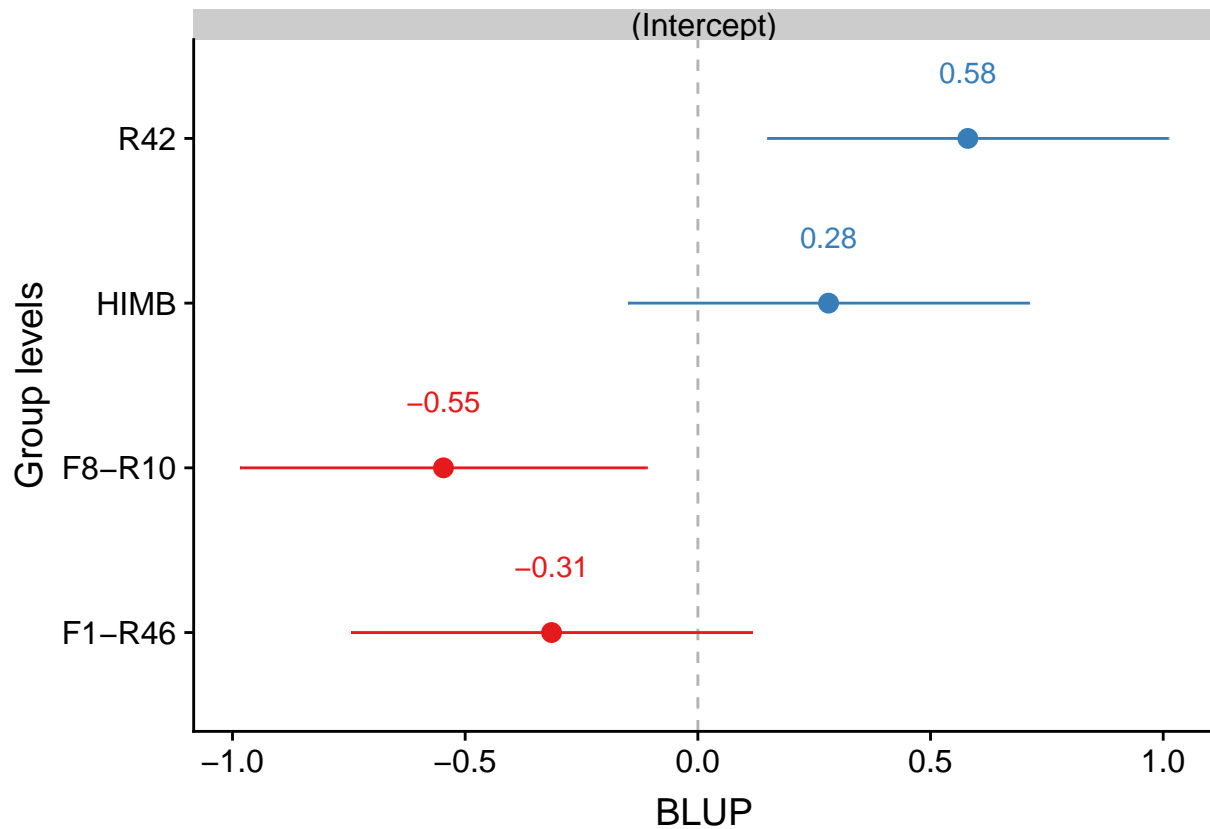
```
## $Location
##      (Intercept)
## F1-R46  -0.3143913
## F8-R10  -0.5466722
## HIMB    0.2808627
## R42     0.5802008
```

```
fixef(full)
```

```
##      (Intercept)      Seasonwinter      Light
##      5.42443464      2.64740866      -0.04057347
##      DomD      Seasonwinter:Light      Seasonwinter:DomD
##      1.02243154      -0.14845501      -3.81871720
##      Light:DomD      Seasonwinter:Light:DomD
##      -0.12052316      0.27682361
```

```
sjp.lmer(full, y.offset = .4)
```





Chlorophyll per symbiont cell

```
##### chlcell --
Y<-model.data$chlcell
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use additive model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df   AIC   BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1    6 13.914 30.588 -0.95686   1.9137
## object 10 18.546 46.337  0.72694  -1.4539 3.3676    4    0.4983
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 22.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.71851 -0.51819 -0.04947  0.47298  2.71238
##
```

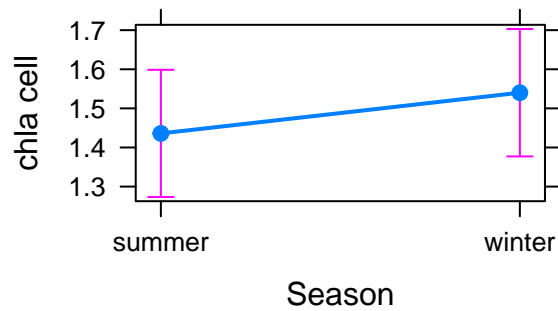
```

## Random effects:
##   Groups   Name      Variance Std.Dev.
##   Location (Intercept) 0.02269  0.1506
##   Residual          0.05655  0.2378
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)   1.818462   0.094137   5.970000  19.317 1.31e-06 ***
## Seasonwinter   0.104004   0.049089  112.760000   2.119  0.0363 *
## Light         -0.028034   0.005004  114.570000  -5.603 1.47e-07 ***
## DomD          -0.546264   0.054293  112.810000 -10.061 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) Ssnwnt Light
## Seasonwintr -0.419
## Light       -0.486  0.456
## DomD        0.089 -0.244 -0.426
print(anova(add, type=2), digits=5)

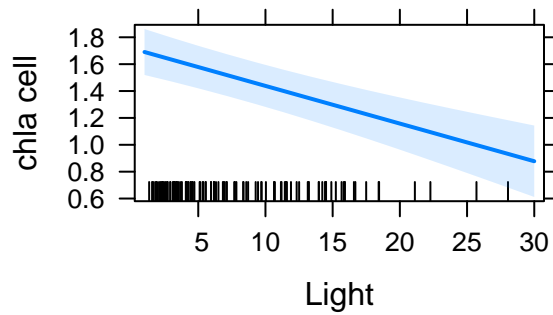
## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##              Sum Sq Mean Sq NumDF DenDF F.value    Pr(>F)
## Season 0.2539  0.2539      1 112.76   4.489  0.03631 *
## Light  1.7754  1.7754      1 114.57  31.393 1.472e-07 ***
## Dom    5.7251  5.7251      1 112.81 101.232 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
plot(allEffects(add), ylab="chla cell", par.strip.text=list(cex=0.7))

```

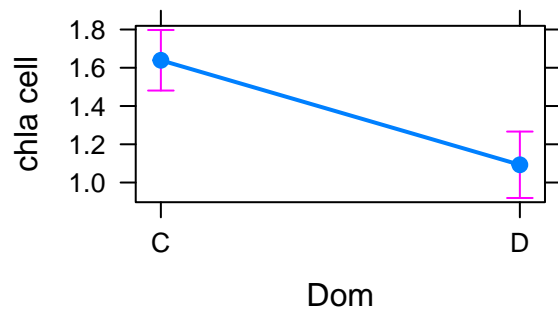
### Season effect plot



### Light effect plot



### Dom effect plot



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46 -0.01688154
## F8-R10 -0.19764101
## HIMB   0.10090019
## R42    0.11362236
```

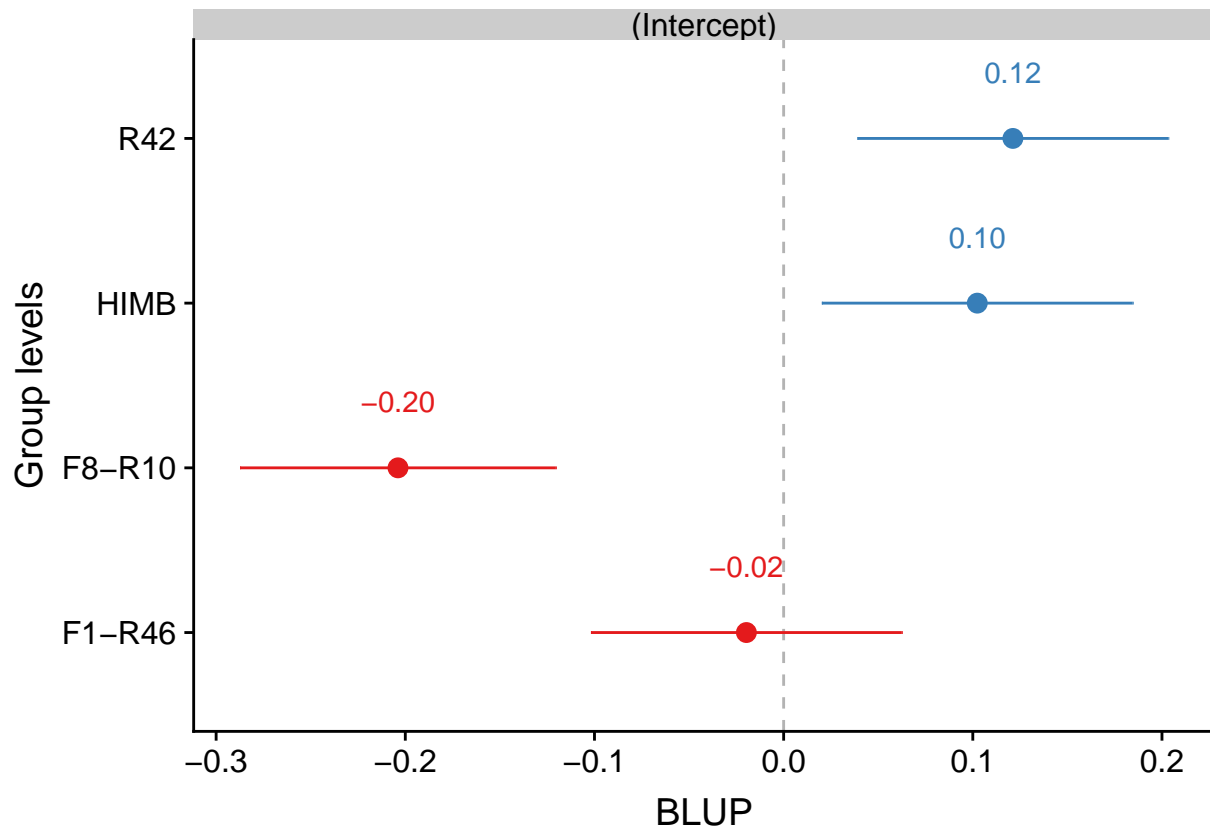
```
rand(add)
```

```
## Analysis of Random effects Table:
##           Chi.sq Chi.DF p.value
## Location   19.8      1 9e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
## (Intercept) Seasonwinter      Light      DomD
##  1.81846170  0.10400380 -0.02803423 -0.54626369
```

```
sjp.lmer(full, y.offset = .4)
```



isotopes

host d13C

```
##### host..d13C --
Y<-model.data$host..d13C
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use additive model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1     6 338.78 355.45 -163.39  326.78
## object 10 343.81 371.60 -161.91  323.81 2.9695    4    0.5629
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 337
##
## Scaled residuals:
```

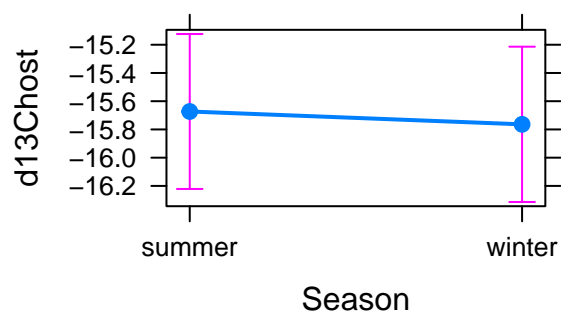
```

##      Min      1Q   Median      3Q      Max
## -2.66981 -0.68614 -0.03878  0.69141  2.21319
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   Location (Intercept) 0.2409   0.4908
##   Residual              0.8772   0.9366
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)  -16.02988    0.33087    7.61000 -48.448 9.33e-11 ***
## Seasonwinter  -0.09107    0.19318  113.10000  -0.471 0.638238
## Light          0.07569    0.01964  114.96000   3.854 0.000192 ***
## DomD           -0.96124    0.21365  113.18000  -4.499 1.66e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) Ssnwnt Light
## Seasonwintr  -0.468
## Light         -0.543  0.455
## DomD           0.099 -0.243 -0.425
print(anova(add, type=2), digits=5)

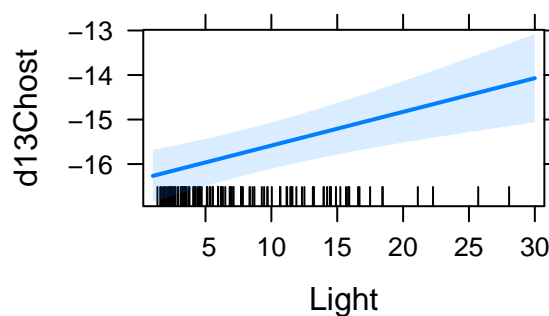
## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##      Sum Sq Mean Sq NumDF  DenDF F.value    Pr(>F)
## Season  0.195    0.195     1 113.10  0.2223 0.6382382
## Light  13.029   13.029     1 114.96 14.8532 0.0001919 ***
## Dom    17.757   17.757     1 113.18 20.2430 1.662e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
plot(allEffects(add), ylab="d13Chost", par.strip.text=list(cex=0.7))

```

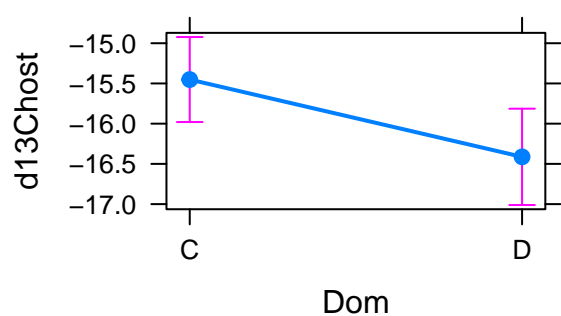
### Season effect plot



### Light effect plot



### Dom effect plot



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46  -0.2856087
## F8-R10   0.3580103
## HIMB    -0.4978117
## R42      0.4254101
```

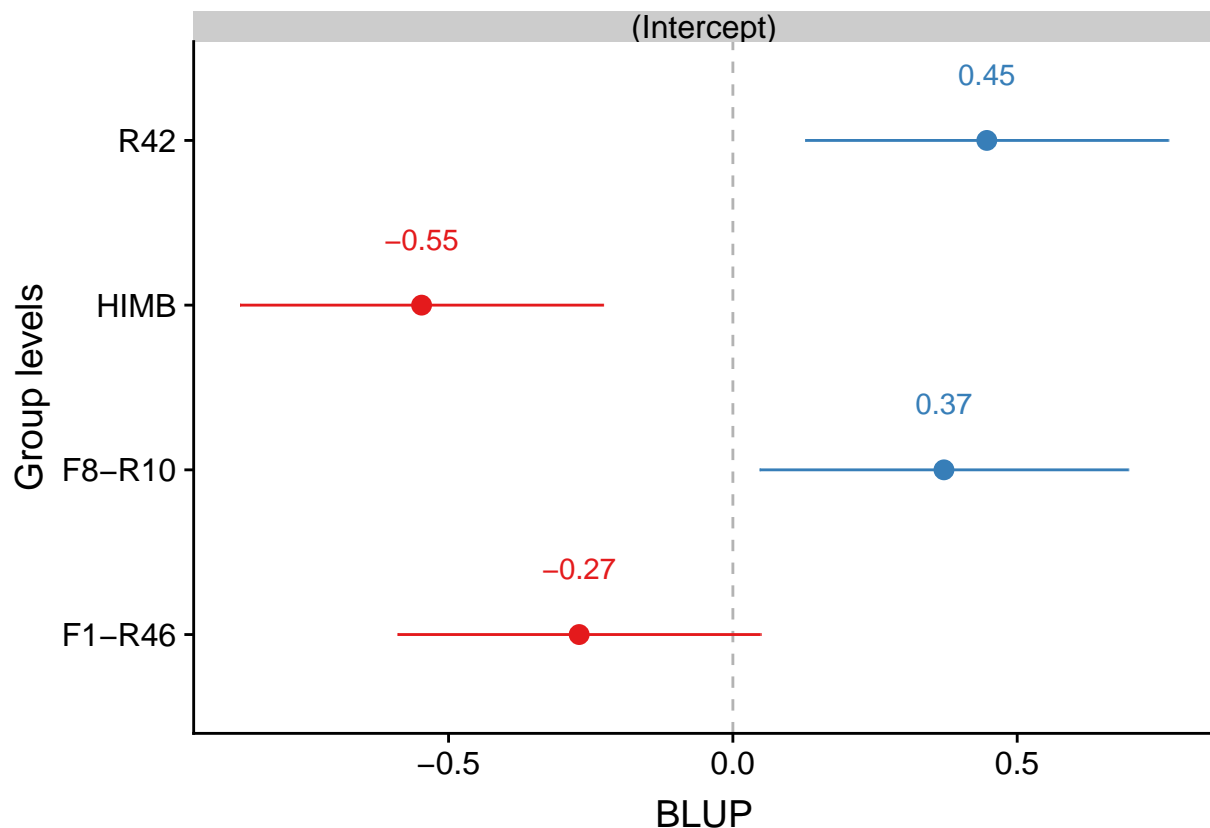
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location  15.4      1 9e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
## (Intercept) Seasonwinter      Light      DomD
## -16.02988136 -0.09107261  0.07569224 -0.96124447
```

```
sjp.lmer(full, y.offset = .4)
```



symbiont d13C

```
##### symb..d13C --
Y<-model.data$symb..d13C
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use additive model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1     6 353.38 370.05 -170.69   341.38
## object 10 354.11 381.90 -167.05   334.11 7.2693     4    0.1223
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 351
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.84081 -0.67259 -0.02418  0.72635  2.47022
##
```

```

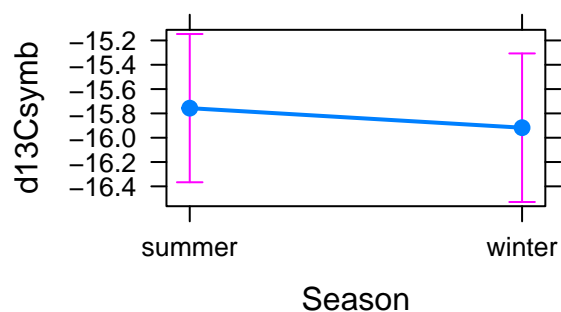
## Random effects:
## Groups Name Variance Std.Dev.
## Location (Intercept) 0.3044 0.5518
## Residual 0.9882 0.9941
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) -16.29310 0.36289 7.08000 -44.899 5.93e-10 ***
## Seasonwinter -0.16092 0.20510 113.00000 -0.785 0.4343
## Light 0.08666 0.02087 114.87000 4.153 6.34e-05 ***
## DomD -0.64063 0.22683 113.07000 -2.824 0.0056 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) Ssnwnt Light
## Seasonwintr -0.453
## Light -0.526 0.455
## DomD 0.096 -0.243 -0.426
print(anova(add, type=2), digits=5)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season 0.6084 0.6084 1 113.00 0.6156 0.434329
## Light 17.0411 17.0411 1 114.87 17.2439 6.344e-05 ***
## Dom 7.8828 7.8828 1 113.07 7.9766 0.005602 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
plot(allEffects(add), ylab="d13Csymb", par.strip.text=list(cex=0.7))

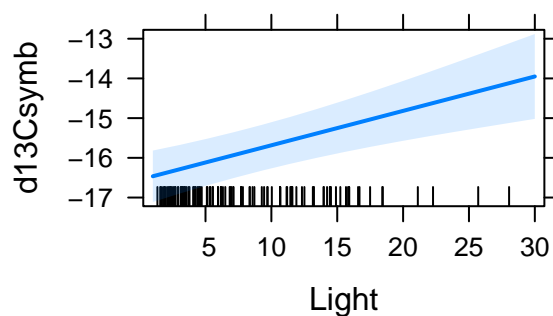
```



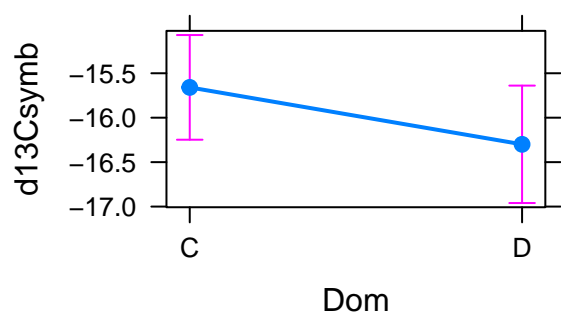
### Season effect plot



### Light effect plot



### Dom effect plot



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46  -0.3459489
## F8-R10   0.5649621
## HIMB    -0.5305526
## R42      0.3115394
```

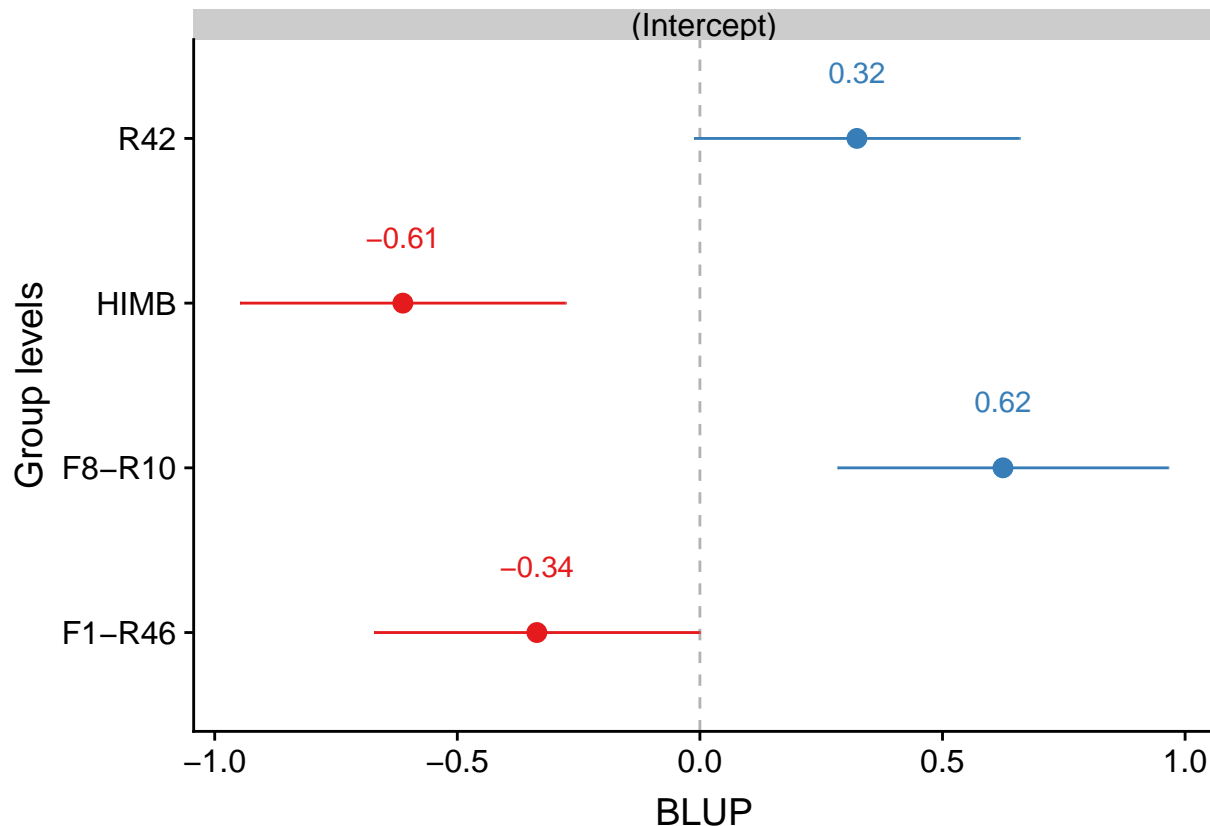
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location  17.2     1 3e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
##      (Intercept) Seasonwinter      Light      DomD
## -16.29310412  -0.16092093   0.08666202 -0.64063417
```

```
sjp.lmer(full, y.offset = .4)
```



host-symbiont d13C

```
##### d13C..host.sym --
Y<-model.data$d13C..host.sym
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ Season:Light +Season:Dom +(1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use add model

## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + Season:Light + Season:Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1    8 27.618 49.851 -5.8091   11.618
## object 10 31.361 59.152 -5.6803   11.361 0.2575    2    0.8792

summary(add)

## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula:
## Y ~ Season + Light + Dom + Season:Light + Season:Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 42.3
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -5.0874 -0.4662  0.0354  0.4346  3.0095
```

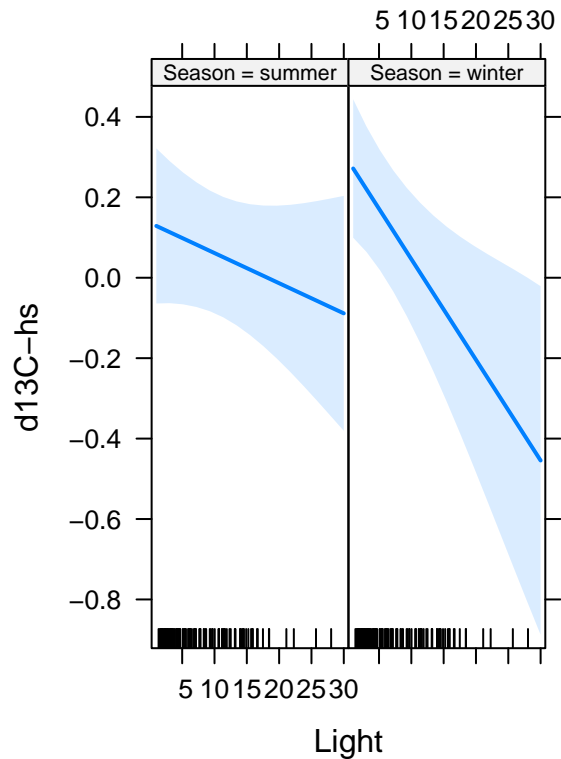
```
##
## Random effects:
## Groups Name Variance Std.Dev.
## Location (Intercept) 0.01849 0.1360
## Residual 0.06304 0.2511
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 0.161776 0.095670 8.780000 1.691 0.12596
## Seasonwinter 0.247105 0.087745 110.010000 2.816 0.00576 **
## Light -0.007499 0.006498 111.750000 -1.154 0.25099
## DomD -0.091643 0.087006 110.550000 -1.053 0.29450
## Seasonwinter:Light -0.017540 0.010292 110.650000 -1.704 0.09115 .
## Seasonwinter:DomD -0.312838 0.115375 110.720000 -2.711 0.00777 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) Ssnwnt Light DomD Ssnw:L
## Seasonwintr -0.513
## Light -0.585 0.592
## DomD 0.128 -0.116 -0.508
## Ssnwntr:Lgh 0.326 -0.757 -0.559 0.285
## Ssnwntr:DmD -0.088 0.023 0.370 -0.753 -0.372

print(anova(add, type=2), digits=4)

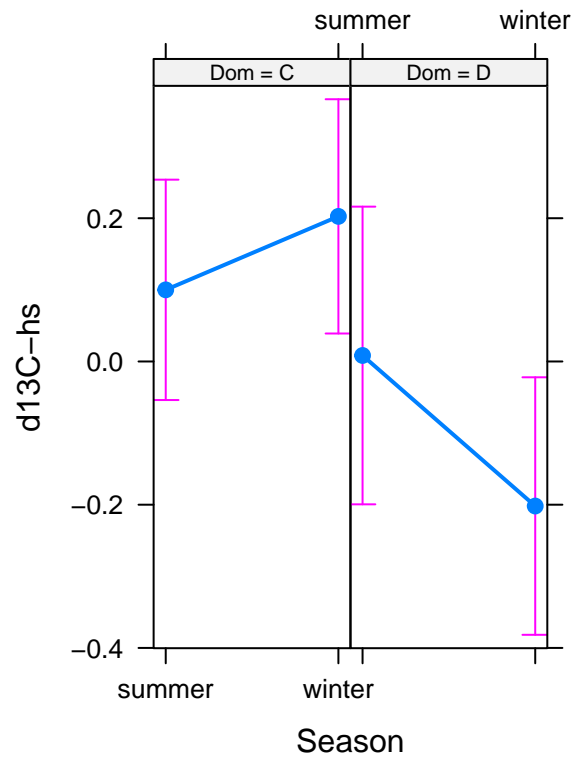
## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season 0.5252 0.5252 1 110.0 8.330 0.00469 **
## Light 0.4347 0.4347 1 113.0 6.896 0.00984 **
## Dom 1.4148 1.4148 1 111.0 22.443 6.45e-06 ***
## Season:Light 0.1831 0.1831 1 110.7 2.904 0.09115 .
## Season:Dom 0.4635 0.4635 1 110.7 7.352 0.00777 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

plot(allEffects(add), ylab="d13C-hs", par.strip.text=list(cex=0.7))
```

### Season\*Light effect plot



### Season\*Dom effect plot



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46  0.04127749
## F8-R10 -0.18899031
## HIMB    0.05229900
## R42     0.09541382
```

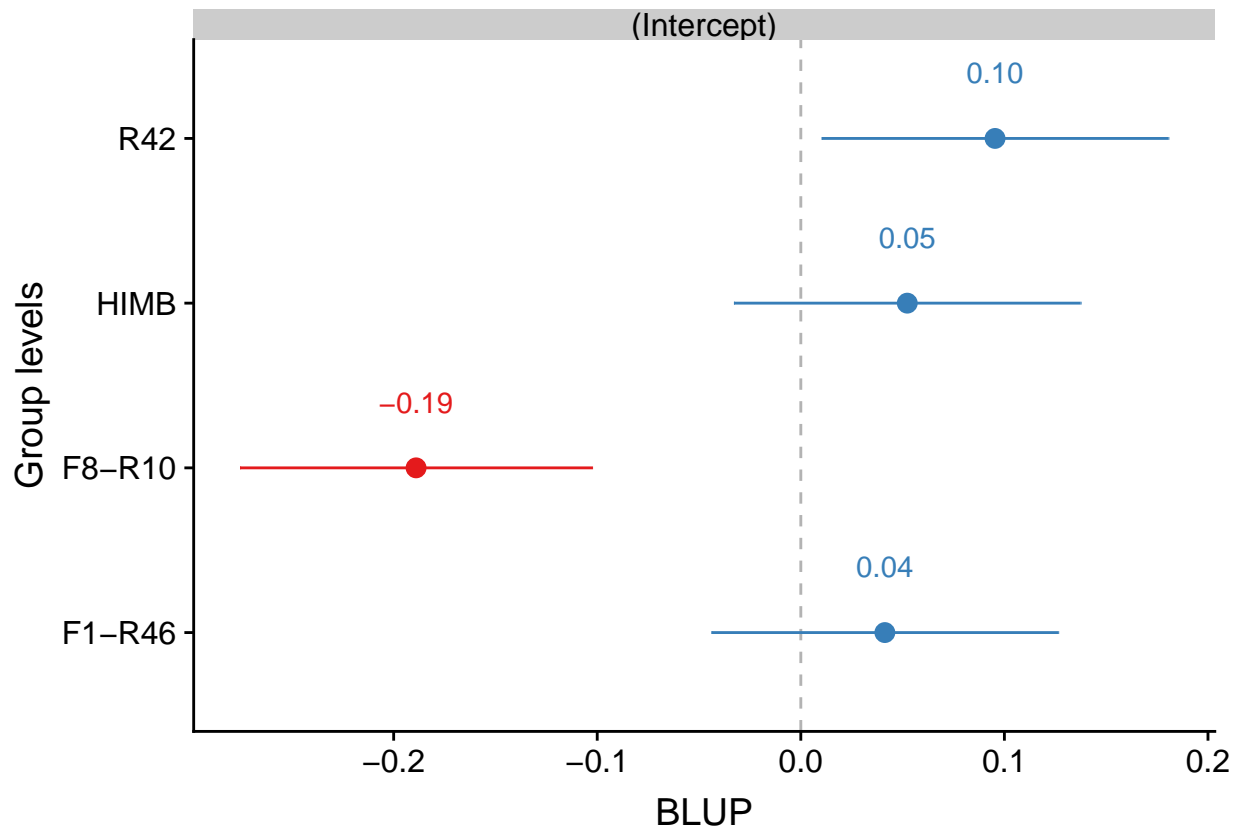
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location    13     1 3e-04 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
##      (Intercept)      Seasonwinter      Light
##      0.16177557      0.24710514      -0.00749862
##      DomD Seasonwinter:Light Seasonwinter:DomD
##      -0.09164343      -0.01754001      -0.31283841
```

```
sjp.lmer(add, y.offset = .4)
```



skeleton d13C

```
##### d13C..skel --
Y<-model.data$d13C..skel
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use full model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1     6 310.41 327.09 -149.21  298.41
## object 10 315.48 343.27 -147.74  295.48 2.9331    4    0.5691
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 309.9
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.96247 -0.80700  0.02393  0.68927  2.44873
##
```

```

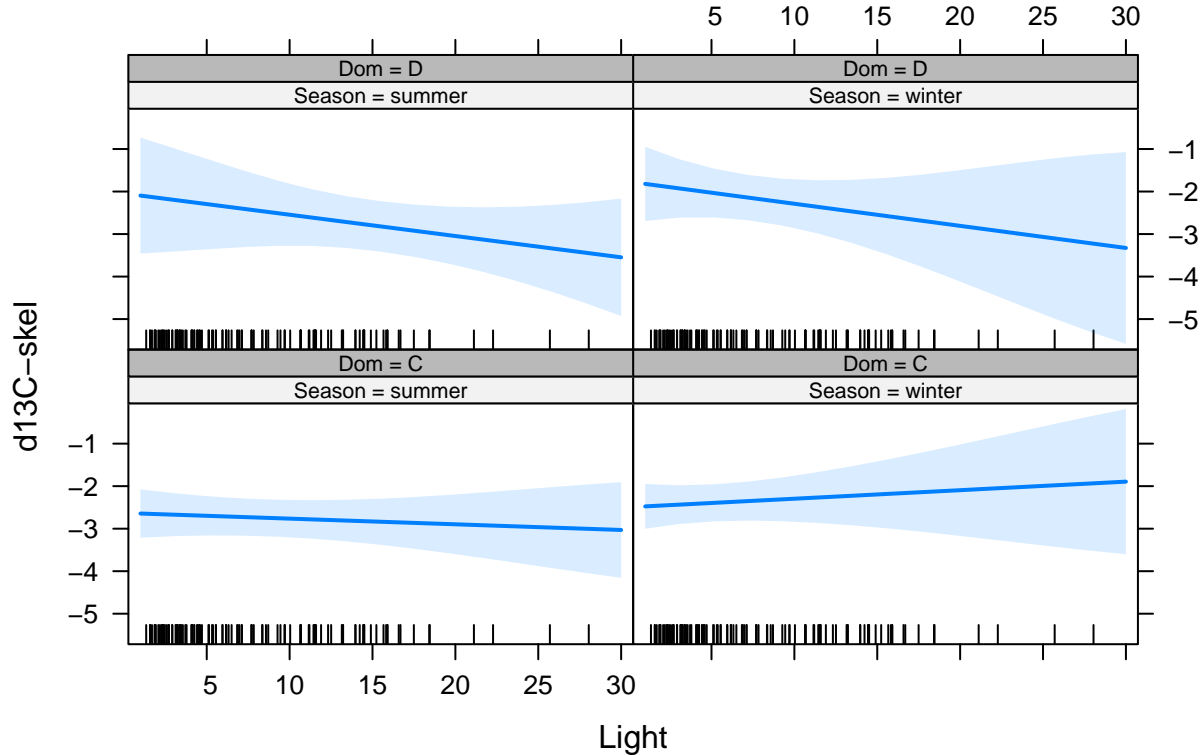
## Random effects:
## Groups Name Variance Std.Dev.
## Location (Intercept) 0.1225 0.3501
## Residual 0.7003 0.8368
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) -2.62727 0.26398 10.18000 -9.953 1.44e-06 ***
## Seasonwinter 0.35828 0.17239 113.52000 2.078 0.0399 *
## Light -0.01686 0.01745 114.75000 -0.966 0.3360
## DomD 0.15238 0.19064 113.62000 0.799 0.4258
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) Ssnwnt Light
## Seasonwintr -0.521
## Light -0.605 0.453
## DomD 0.108 -0.241 -0.423
print(anova(add, type=2), digits=4)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season 3.0248 3.0248 1 113.5 4.319 0.0399 *
## Light 0.6538 0.6538 1 114.8 0.934 0.3360
## Dom 0.4474 0.4474 1 113.6 0.639 0.4258
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
posthoc<-emmeans(add, ~Season)
CLD(posthoc, Letters=letters)

## Season emmean SE df lower.CL upper.CL .group
## summer -2.690017 0.2179220 4.88 -3.254270 -2.125764 a
## winter -2.331733 0.2101285 4.27 -2.901077 -1.762390 b
##
## Results are averaged over the levels of: Dom
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
## significance level used: alpha = 0.05
plot(allEffects(full), ylab="d13C-skel", par.strip.text=list(cex=0.7))

```

## Season\*Light\*Dom effect plot



host d15N

```
##### host..d15N --
Y<-model.data$host..d15N
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+Season:Light+ Season:Dom+ Light:Dom +(1|Location), data=model.data, na.act
anova(full, add) #use add model

## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + Season:Light + Season:Dom + Light:Dom +
## ..1: (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1     9 85.846 110.86 -33.923   67.846
## object 10 87.496 115.29 -33.748   67.496  0.35     1    0.5541

summary(add)

## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula:
## Y ~ Season + Light + Dom + Season:Light + Season:Dom + Light:Dom +
## (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 100.6
##
## Scaled residuals:
```

```

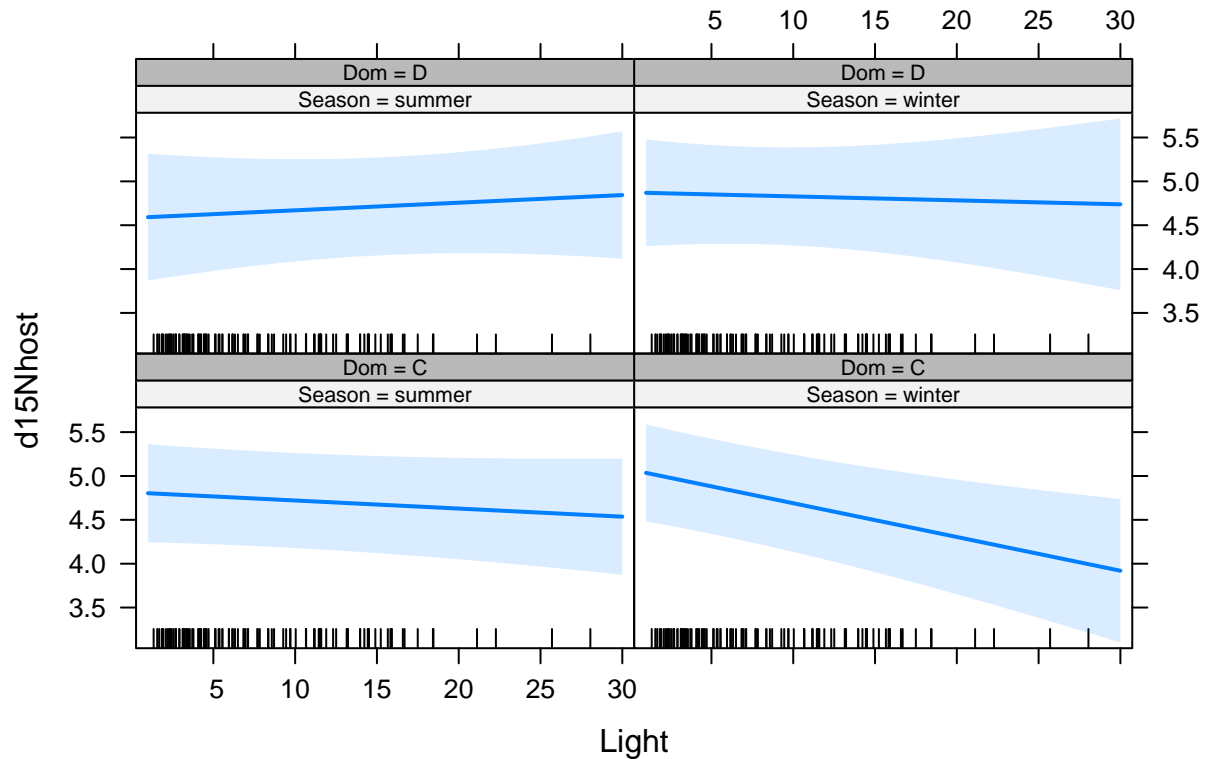
##      Min      1Q   Median      3Q      Max
## -2.87510 -0.49317 -0.02088  0.60805  2.07331
##
## Random effects:
##   Groups   Name                Variance Std.Dev.
##   Location (Intercept) 0.29361  0.5419
##   Residual              0.09454  0.3075
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.825123   0.284627   3.570000  16.952 0.000153 ***
## Seasonwinter    0.232588   0.107773  109.020000   2.158 0.033113 *
## Light          -0.010659   0.008625  109.160000  -1.236 0.219161
## DomD           -0.321879   0.218258  109.080000  -1.475 0.143157
## Seasonwinter:Light -0.024721  0.012630  109.080000  -1.957 0.052870 .
## Seasonwinter:DomD  0.187905   0.167453  109.070000   1.122 0.264268
## Light:DomD       0.024532   0.013811  109.050000   1.776 0.078484 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) Ssnwnt Light  DomD   Ssnw:L Ssn:DD
## Seasonwintr -0.218
## Light        -0.261  0.572
## DomD          -0.059  0.009  0.097
## Ssnwntr:Lgh  0.130 -0.752 -0.498  0.167
## Ssnwntr:DmD  0.021 -0.021  0.089 -0.777 -0.332
## Light:DomD   0.098 -0.075 -0.376 -0.872 -0.034  0.535
print(anova(add, type=2), digits=4)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##              Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season        0.4478  0.4478     1 109.0   4.737 0.0317 *
## Light          0.3877  0.3877     1 109.5   4.101 0.0453 *
## Dom            0.0847  0.0847     1 109.1   0.896 0.3461
## Season:Light  0.3622  0.3622     1 109.1   3.831 0.0529 .
## Season:Dom    0.1190  0.1190     1 109.1   1.259 0.2643
## Light:Dom     0.2983  0.2983     1 109.0   3.155 0.0785 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
plot(allEffects(full), ylab="d15Nhost", par.strip.text=list(cex=0.7))

```



## Season\*Light\*Dom effect plot



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46  0.37281267
## F8-R10  0.05415004
## HIMB    0.35101369
## R42     -0.77797639
```

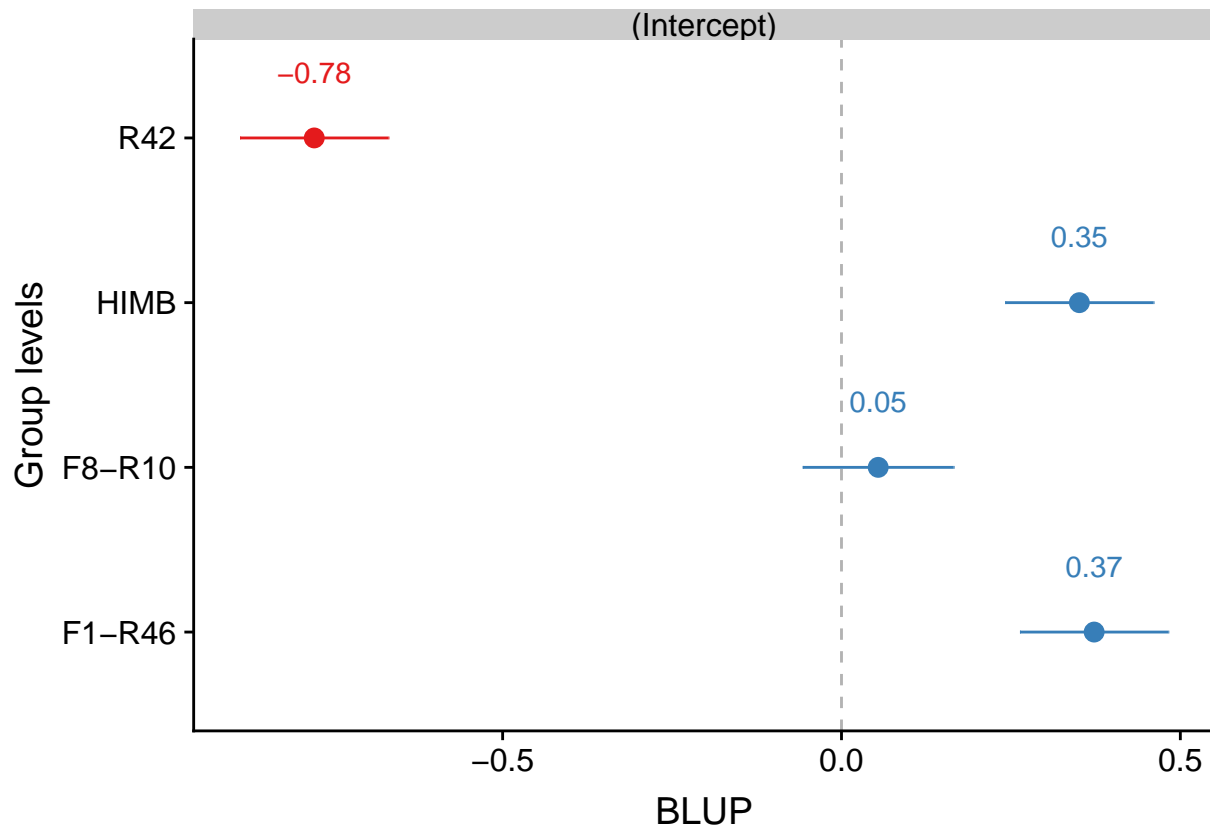
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location    121      1 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
##      (Intercept)      Seasonwinter      Light
##      4.82512285      0.23258755      -0.01065915
##      DomD Seasonwinter:Light Seasonwinter:DomD
##      -0.32187911      -0.02472116      0.18790533
##      Light:DomD
##      0.02453174
```

```
sjp.lmer(add, y.offset = .4)
```



symbiont d15N

```
##### symb..d15N --
Y<-model.data$symb..d15N
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+Season:Light+ Season:Dom+ Light:Dom + (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use full model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + Season:Light + Season:Dom + Light:Dom +
## ..1: (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1    9 94.163 119.17 -38.082   76.163
## object 10 95.262 123.05 -37.631   75.262 0.9013    1    0.3424
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula:
## Y ~ Season + Light + Dom + Season:Light + Season:Dom + Light:Dom +
## (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 108.1
```

```

##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.1583 -0.6369  0.1293  0.6237  2.2340
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   Location (Intercept) 0.4455   0.6675
##   Residual          0.1002   0.3165
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.258107   0.345609    3.390000  12.321 0.000618 ***
## Seasonwinter    0.017130   0.110941  109.010000   0.154 0.877575
## Light           0.005542   0.008879  109.110000   0.624 0.533867
## DomD            -0.453814   0.224684  109.060000  -2.020 0.045857 *
## Seasonwinter:Light -0.011895  0.013002  109.050000  -0.915 0.362297
## Seasonwinter:DomD  0.548332   0.172381  109.050000   3.181 0.001912 **
## Light:DomD       0.026723   0.014217  109.040000   1.880 0.062838 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) Ssnwntr Light  DomD   Ssnw:L Ssn:DD
## Seasonwintr -0.185
## Light        -0.221  0.572
## DomD          -0.050  0.009  0.097
## Ssnwntr:Lgh  0.110 -0.752 -0.498  0.167
## Ssnwntr:DmD  0.018 -0.021  0.089 -0.777 -0.332
## Light:DomD   0.083 -0.075 -0.376 -0.872 -0.033  0.534
print(anova(add, type=2), digits=3)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##              Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season        0.004   0.004     1    109   0.04 0.8377
## Light         0.140   0.140     1    109   1.39 0.2404
## Dom           0.056   0.056     1    109   0.56 0.4552
## Season:Light  0.084   0.084     1    109   0.84 0.3623
## Season:Dom    1.014   1.014     1    109  10.12 0.0019 **
## Light:Dom     0.354   0.354     1    109   3.53 0.0628 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

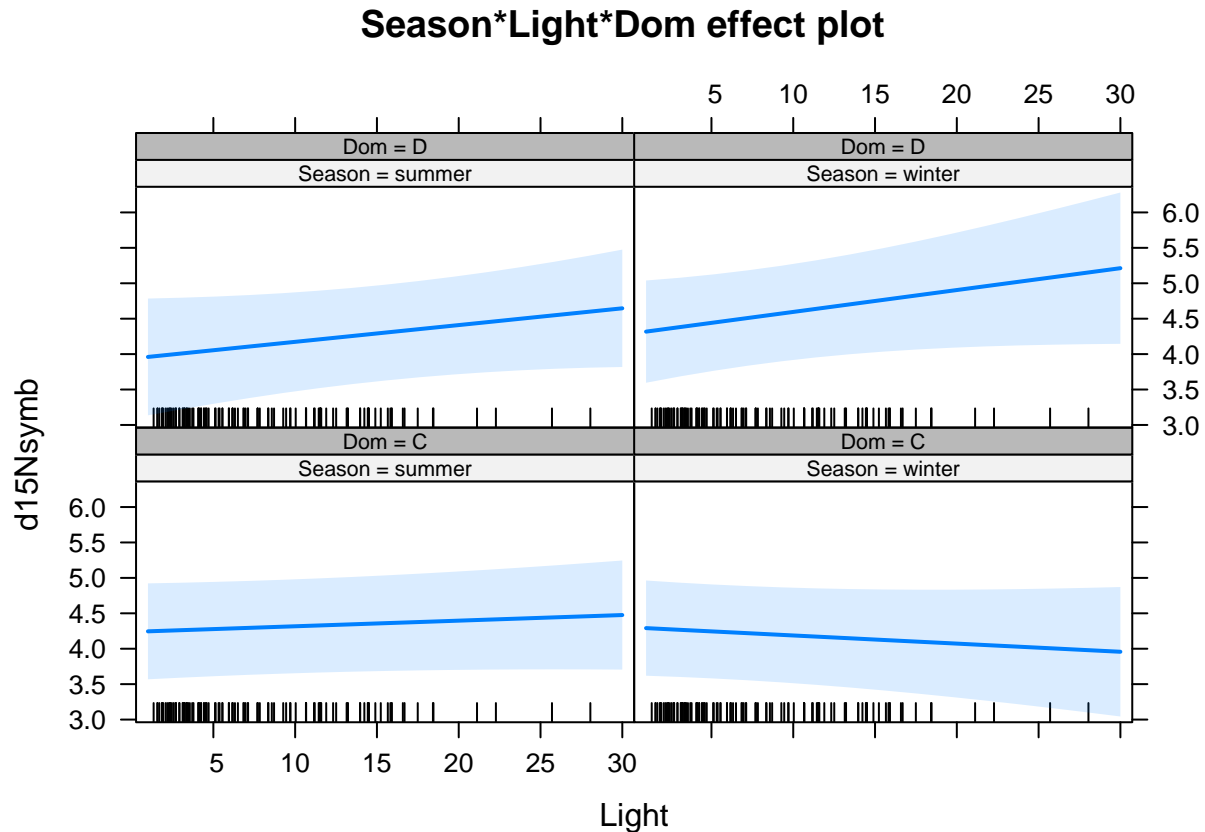
posthoc<-emmeans(full, ~Season*Dom)
CLD(posthoc, Letters=letters)

##   Season Dom   emmean      SE    df lower.CL upper.CL .group
##   summer D   4.131481 0.3610092 4.17 3.144705 5.118257   ab
##   winter C   4.207585 0.3368824 3.16 3.166021 5.249150    a
##   summer C   4.302891 0.3343979 3.07 3.252439 5.353343    a
##   winter D   4.540954 0.3396607 3.27 3.508523 5.573385    b
##

```

```
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

```
plot(allEffects(full), ylab="d15Nsymb", par.strip.text=list(cex=0.7))
```



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46    0.4132212
## F8-R10    0.1547304
## HIMB      0.4122555
## R42       -0.9802071
```

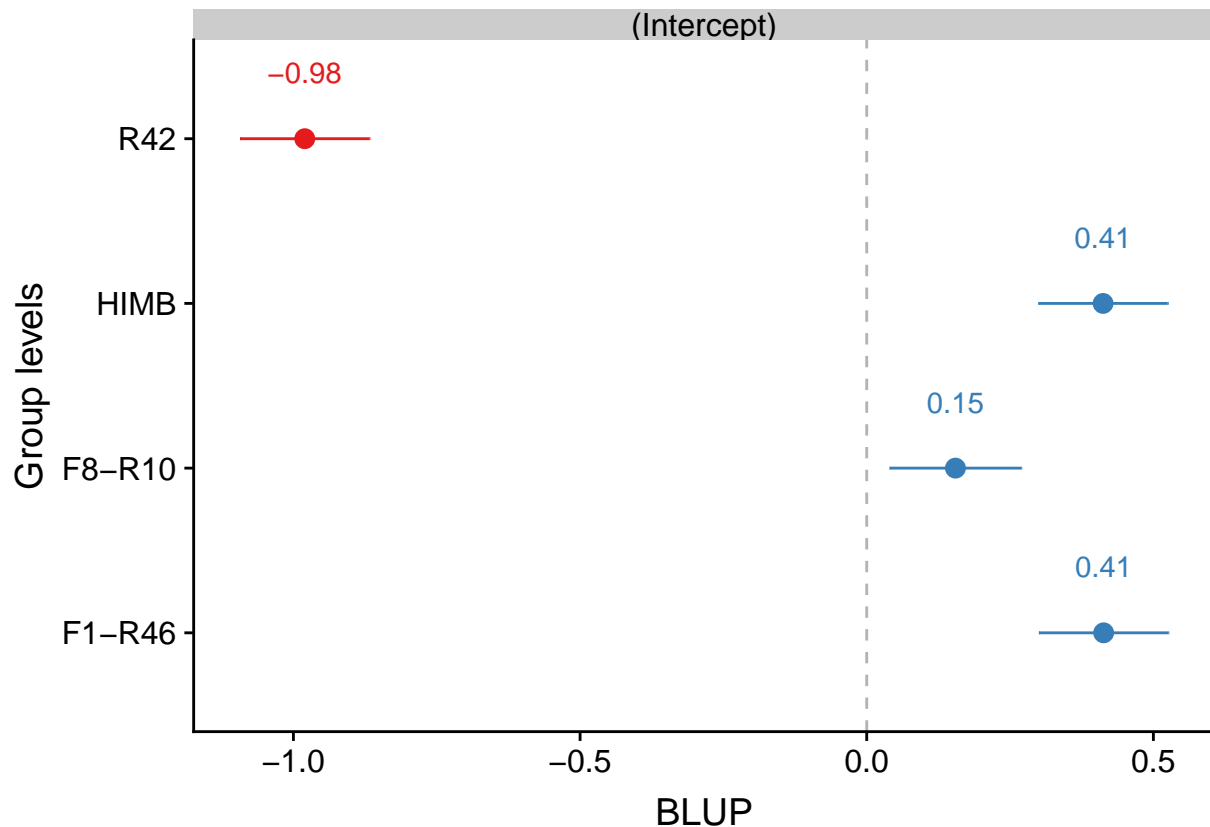
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location    148      1 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
##      (Intercept)      Seasonwinter      Light
##      4.258107245      0.017130046      0.005541514
##      DomD Seasonwinter:Light Seasonwinter:DomD
##      -0.453814481      -0.011894857      0.548331654
```

```
##      Light:DomD
##      0.026722563
sjp.lmer(add, y.offset = .4)
```



host-symbiont d15N

```
##### d15N..host.sym --
Y<-model.data$d15N..host.sym
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ Season:Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use full model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + Season:Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1      7 -7.7053 11.748 10.853 -21.705
## object 10 -4.1556 23.636 12.078 -24.156 2.4503      3      0.4843
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + Season:Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: 3
##
```

```
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.87449 -0.73818  0.03548  0.69225  2.33677
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   Location (Intercept) 0.01553  0.1246
##   Residual          0.04706  0.2169
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    0.601672   0.080872   6.670000    7.440 0.000183 ***
## Seasonwinter    0.135546   0.049510  111.600000    2.738 0.007201 **
## Light          -0.020248   0.004662  113.850000   -4.343 3.06e-05 ***
## DomD            0.126465   0.072076  111.850000    1.755 0.082064 .
## Seasonwinter:DomD -0.399543   0.092535  111.800000   -4.318 3.42e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) Ssnwnt Light  DomD
## Seasonwintr -0.417
## Light        -0.496  0.312
## DomD          0.038  0.159 -0.439
## Ssnwntr:DmD  0.036 -0.427  0.211 -0.727

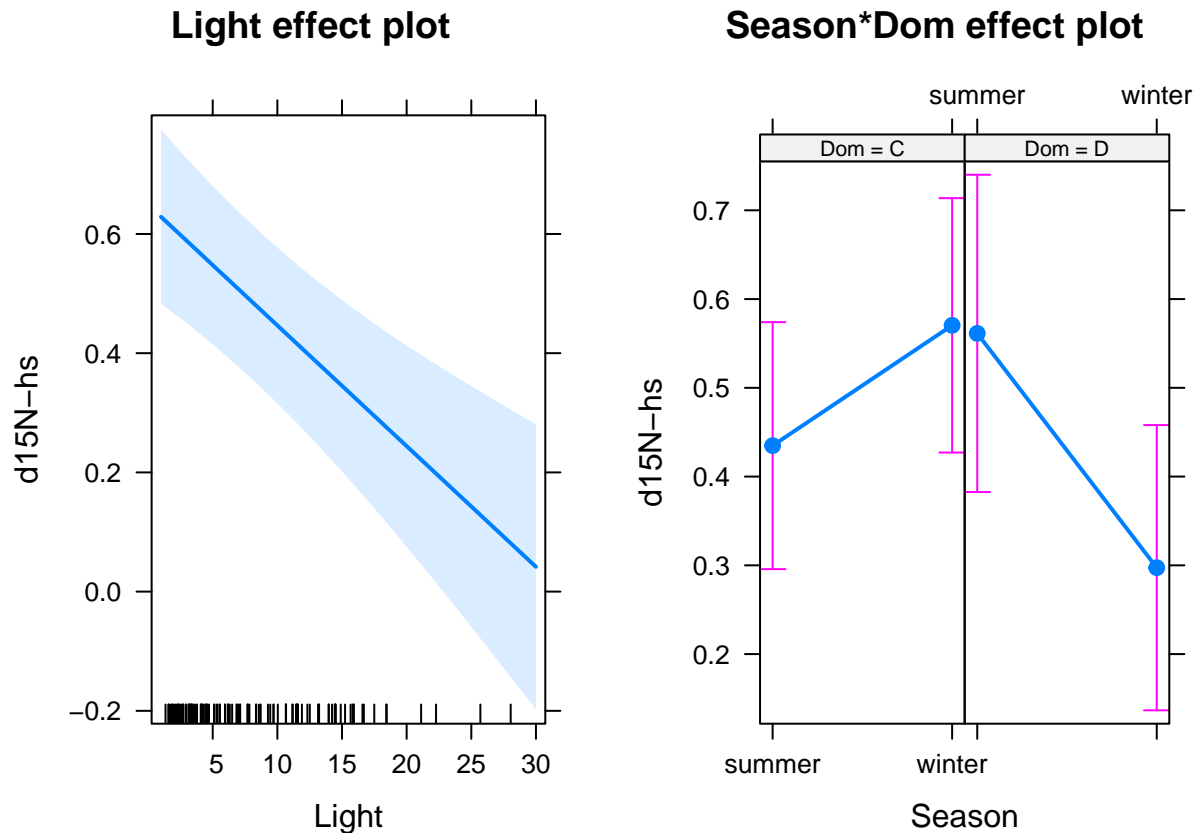
print(anova(add, type=2), digits=3)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##              Sum Sq Mean Sq NumDF DenDF F.value  Pr(>F)
## Season        0.040   0.040     1   112     0.85   0.358
## Light         0.888   0.888     1   114    18.86 3.1e-05 ***
## Dom           0.194   0.194     1   112     4.13   0.044 *
## Season:Dom    0.877   0.877     1   112    18.64 3.4e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

posthoc<-emmeans(full, ~Season*Dom)
CLD(posthoc, Letters=letters)

##   Season Dom   emmean      SE    df  lower.CL  upper.CL .group
## winter D    0.2936771 0.08195540  6.80 0.09871493 0.4886394  a
## summer C    0.4335578 0.07100439  3.88 0.23408224 0.6330333  ab
## summer D    0.5112646 0.11739577 25.04 0.26950469 0.7530245  ab
## winter C    0.5553393 0.07636443  5.13 0.36050544 0.7501731  b
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
## P value adjustment: tukey method for comparing a family of 4 estimates
## significance level used: alpha = 0.05
```

```
plot(allEffects(add), ylab="d15N-hs", par.strip.text=list(cex=0.7))
```



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46 -0.04191457
## F8-R10 -0.08161716
## HIMB   -0.05201146
## R42     0.17554320
```

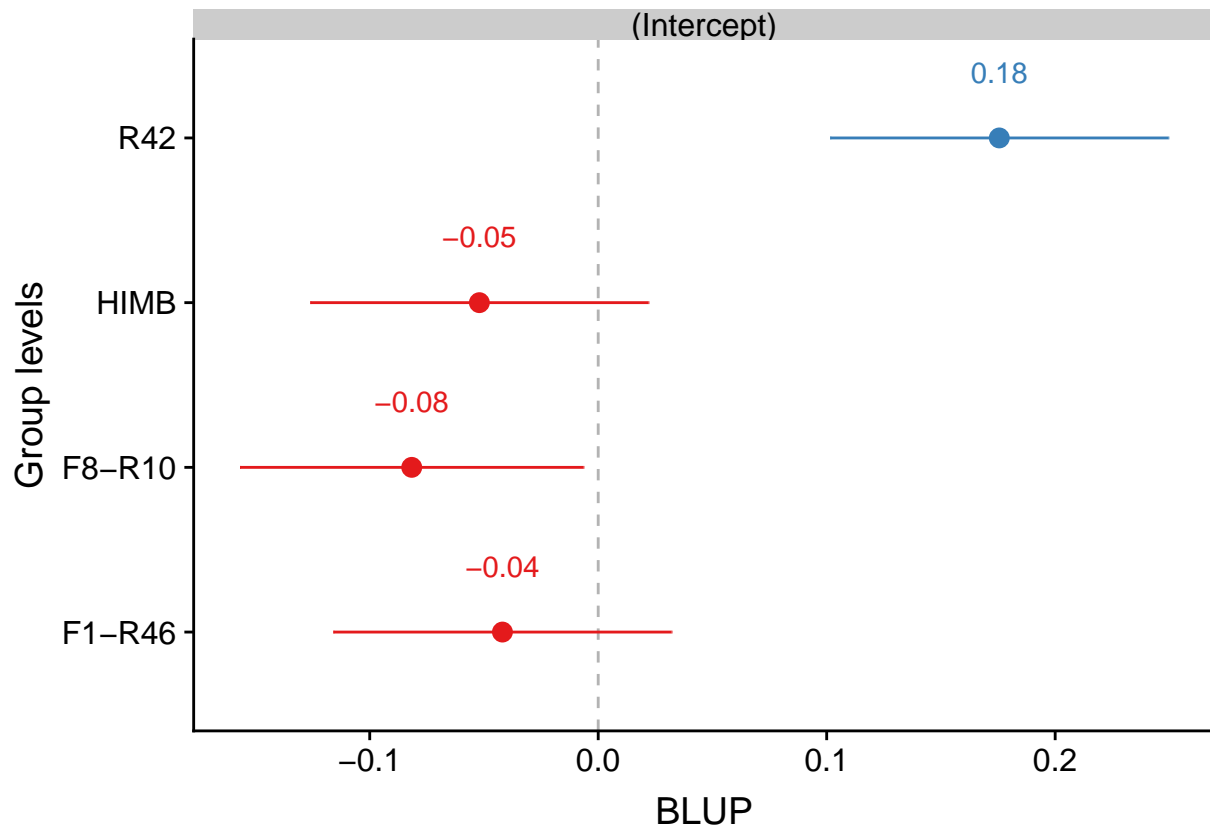
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location  16.4      1 5e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
##      (Intercept)      Seasonwinter      Light      DomD
##      0.60167221      0.13554584     -0.02024803      0.12646507
## Seasonwinter:DomD
##      -0.39954308
```

```
sjp.lmer(add, y.offset = .4)
```



host C:N

```
##### host..C.N --
Y<-model.data$host..C.N
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use full model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1    6 -271.85 -255.18 141.93  -283.85
## object 10 -268.06 -240.27 144.03  -288.06 4.2107    4    0.3782
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: -252.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
```



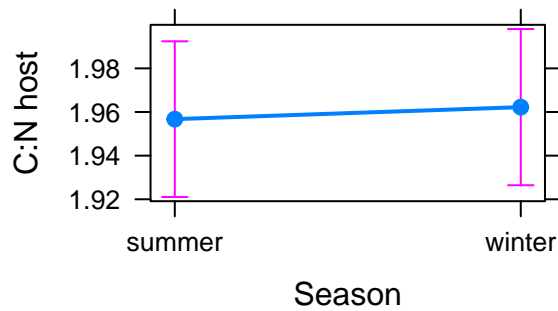
```

## -1.84607 -0.73952 -0.04315 0.66611 3.15900
##
## Random effects:
## Groups Name Variance Std.Dev.
## Location (Intercept) 0.0009024 0.03004
## Residual 0.0052547 0.07249
## Number of obs: 119, groups: Location, 4
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 1.943e+00 2.277e-02 9.890e+00 85.327 1.78e-15 ***
## Seasonwinter 5.501e-03 1.493e-02 1.135e+02 0.368 0.713
## Light 1.382e-03 1.512e-03 1.147e+02 0.914 0.362
## DomD 8.612e-03 1.651e-02 1.136e+02 0.522 0.603
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) Ssnwnt Light
## Seasonwintr -0.523
## Light -0.607 0.452
## DomD 0.108 -0.241 -0.423
print(anova(add, type=2), digits=4)

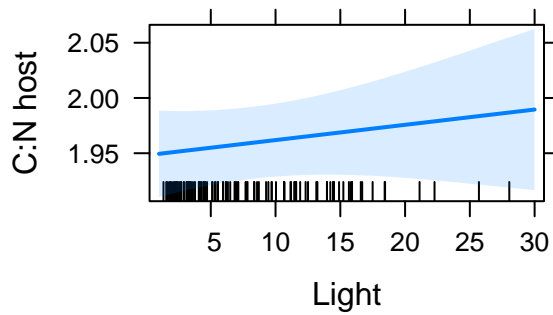
## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Season 0.000713 0.000713 1 113.5 0.1357 0.713
## Light 0.004394 0.004394 1 114.7 0.8361 0.362
## Dom 0.001429 0.001429 1 113.6 0.2720 0.603
plot(allEffects(add), ylab="C:N host", par.strip.text=list(cex=0.7))

```

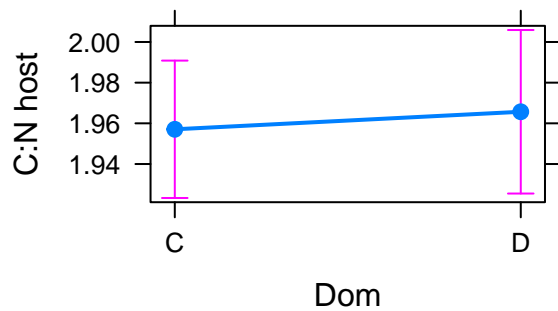
### Season effect plot



### Light effect plot



### Dom effect plot



```
ranef(add)
```

```
## $Location
##      (Intercept)
## F1-R46 -0.00994385
## F8-R10  0.02064641
## HIMB    0.02340875
## R42     -0.03411131
```

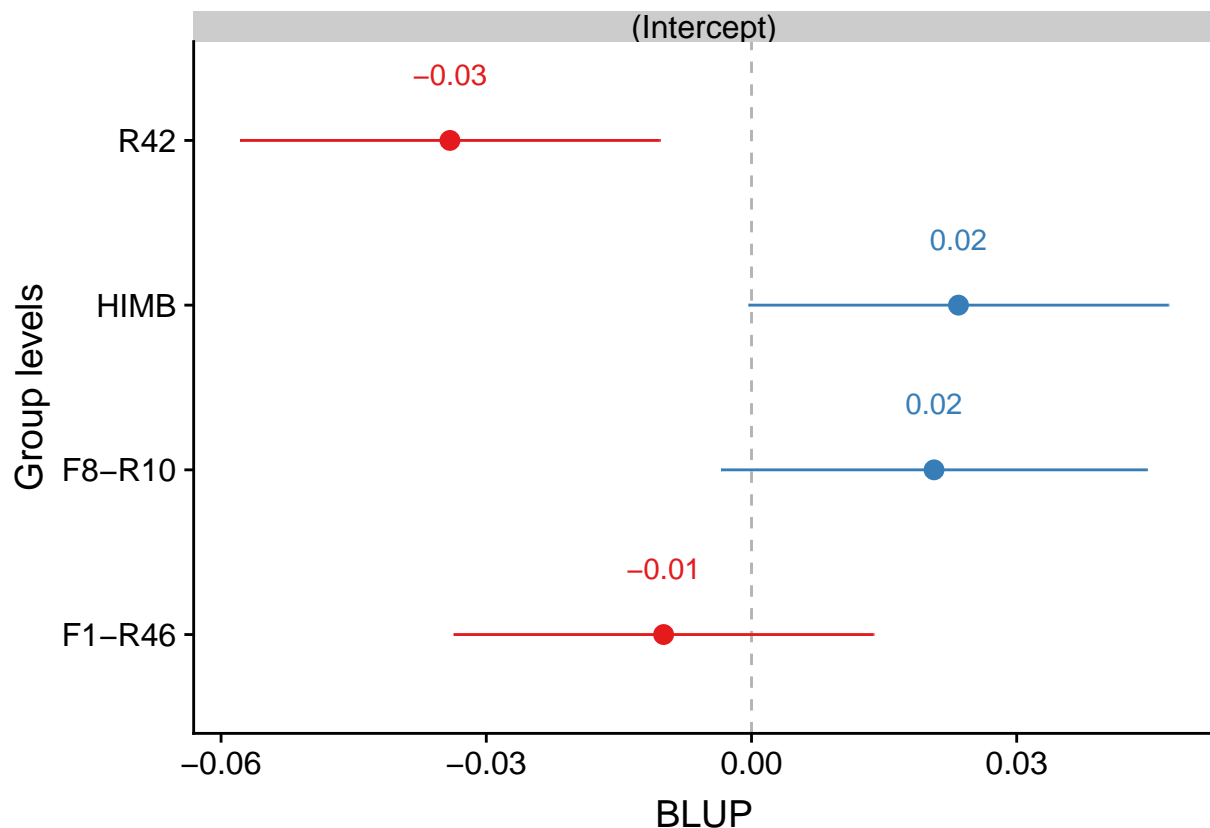
```
rand(add)
```

```
## Analysis of Random effects Table:
##      Chi.sq Chi.DF p.value
## Location   7.57     1  0.006 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fixef(add)
```

```
## (Intercept) Seasonwinter      Light      DomD
## 1.942951666  0.005501474  0.001382129  0.008612347
```

```
sjp.lmer(add, y.offset = .4)
```



symbiont C:N

```
##### symb..C.N --
Y<-model.data$symb..C.N
full<-lmer(Y~Season*Light*Dom+ (1|Location), data=model.data, na.action=na.exclude)
add<-lmer(Y~Season+Light+Dom+ (1|Location), data=model.data, na.action=na.exclude)
anova(full, add) #use full model
```

```
## Data: model.data
## Models:
## ..1: Y ~ Season + Light + Dom + (1 | Location)
## object: Y ~ Season * Light * Dom + (1 | Location)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## ..1    6 -210.70 -194.07 111.35 -222.70
## object 10 -207.75 -180.04 113.88 -227.75 5.0519    4    0.282
```

```
summary(add)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Season + Light + Dom + (1 | Location)
## Data: model.data
##
## REML criterion at convergence: -192
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.8765 -0.6378 -0.1564  0.6934  2.9317
##
```

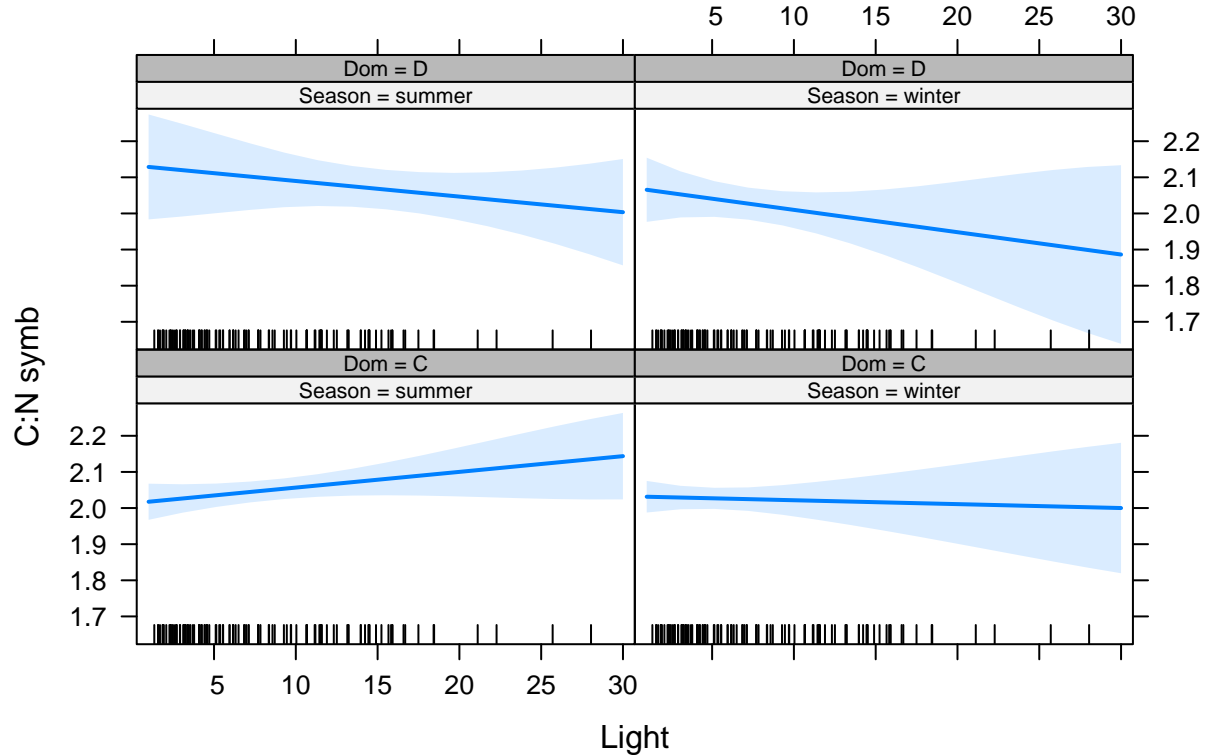
```

## Random effects:
## Groups   Name      Variance Std.Dev.
## Location (Intercept) 0.00000 0.00000
## Residual      0.00918 0.09581
## Number of obs: 118, groups: Location, 4
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)  2.048e+00  2.155e-02  1.140e+02  95.038  <2e-16 ***
## Seasonwinter -2.715e-02  1.945e-02  1.140e+02  -1.396   0.165
## Light        5.587e-04  1.849e-03  1.140e+02   0.302   0.763
## DomD         2.871e-03  2.143e-02  1.140e+02   0.134   0.894
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) Ssnwnt Light
## Seasonwintr  -0.678
## Light        -0.786  0.416
## DomD         0.099 -0.220 -0.394
print(anova(add, type=2), digits=5)

## Analysis of Variance Table of type II with Satterthwaite
## approximation for degrees of freedom
##              Sum Sq   Mean Sq NumDF DenDF F.value Pr(>F)
## Season 0.0178944 0.0178944     1    114  1.94918 0.1654
## Light  0.0008386 0.0008386     1    114  0.09135 0.7630
## Dom    0.0001649 0.0001649     1    114  0.01796 0.8936
plot(allEffects(full), ylab="C:N symb", par.strip.text=list(cex=0.7))

```

## Season\*Light\*Dom effect plot



### Figures

#### qPCR figures

```
## Figures
#####
#####
#####

#qPCR and symbionts

#####
##### Plot Dominant Symbiont and Depth (both seasons)
Symb<-model.data
Symb$Dominant <- ifelse(Symb$Dom=="D", 1, 0)
results.all=glm(Dominant~newDepth, family = "binomial", data = Symb)
anova(results.all, test = "Chisq")
summary(results.all)
fitted.all <- predict(results.all, newdata = list(newDepth=seq(0,12,0.1)), type = "response")
# plot(fitted.all, ylab="proportion D", ylim=c(0,1))

##### summer only symbionts
sum.dat<-Symb[(Symb$Season=="summer"),]
results.sum=glm(Dominant~newDepth, family = "binomial", data = sum.dat)
anova(results.sum, test = "Chisq")
summary(results.sum)
fitted.sum <- predict(results.sum, newdata = list(newDepth=seq(0,12,0.1)), type = "response")
```

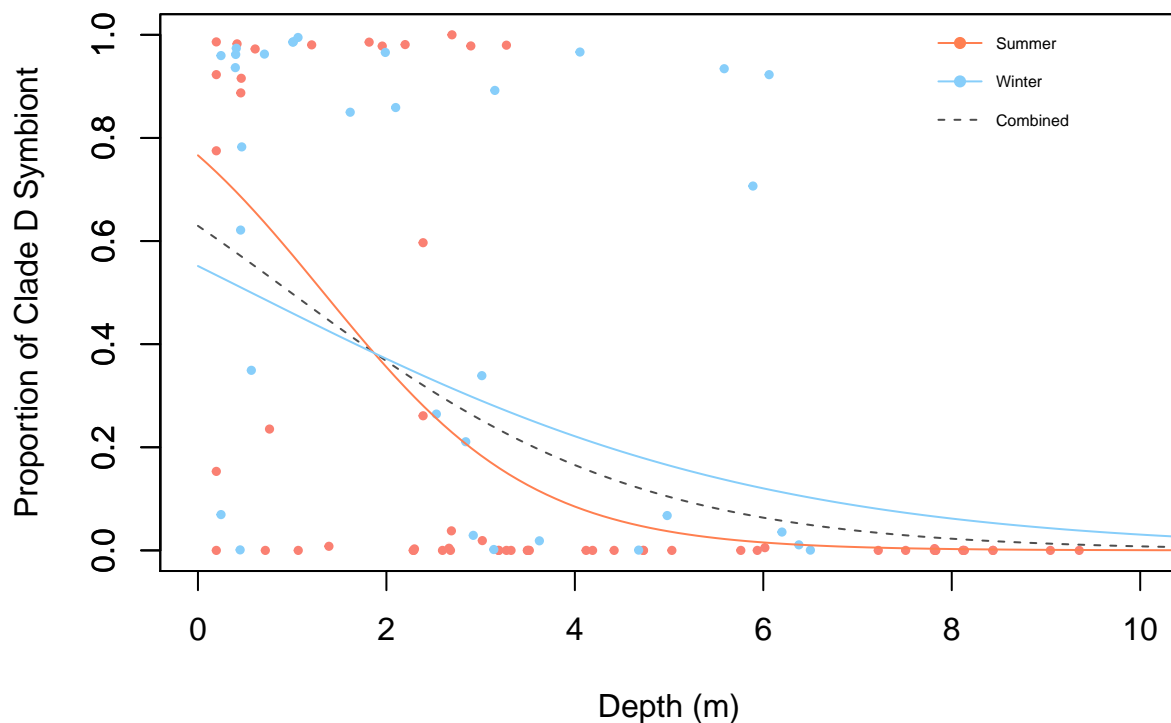
```

# plot(fitted.sum, ylab="proportion D", ylim=c(0,1))

##### winter only symbionts
wint.dat<-Symb[(Symb$Season=="winter"),]
results.win=glm(Dominant~newDepth, family = "binomial", data = wint.dat)
anova(results.win, test = "Chisq")
summary(results.win)
fitted.win <- predict(results.win, newdata = list(newDepth=seq(0,12,0.1)), type = "response")
# plot(fitted.win, ylab="proportion D", ylim=c(0,1))

#####
#####
## Figure of Dominant symbiont clades across seasons
## **Note** where points equal 0.0 is 100% D, where they equal 1.0 is 100% C
par(mar=c(5,4,3,2))
plot(sum.dat$propD~sum.dat$newDepth, xlab="Depth (m)", ylab = "Proportion of Clade D Symbiont", pch=19,
par(new=T)
plot(wint.dat$propD~wint.dat$newDepth, pch=19, col="lightskyblue", xlim=c(0,10), ylim=c(0.0, 1.0), xaxt="n", yaxt="n")
lines(fitted.all ~ seq(0,12,0.1), col="gray30", lwd=1, lty=2)
lines(fitted.sum ~ seq(0,12,0.1), col="coral", lwd=1)
lines(fitted.win ~ seq(0,12,0.1), col="lightskyblue", lwd=1)
legend("topright", pch=c(19,19, NA), lty=c(1,1,2), col=c("coral", "lightskyblue", "gray30"), legend=c("Summer", "Winter", "Combined"))

```



```

dev.copy(pdf, "figures/symbionts/Symbionts_by_Season.pdf", encod="MacRoman", height=4, width=4)
dev.off()

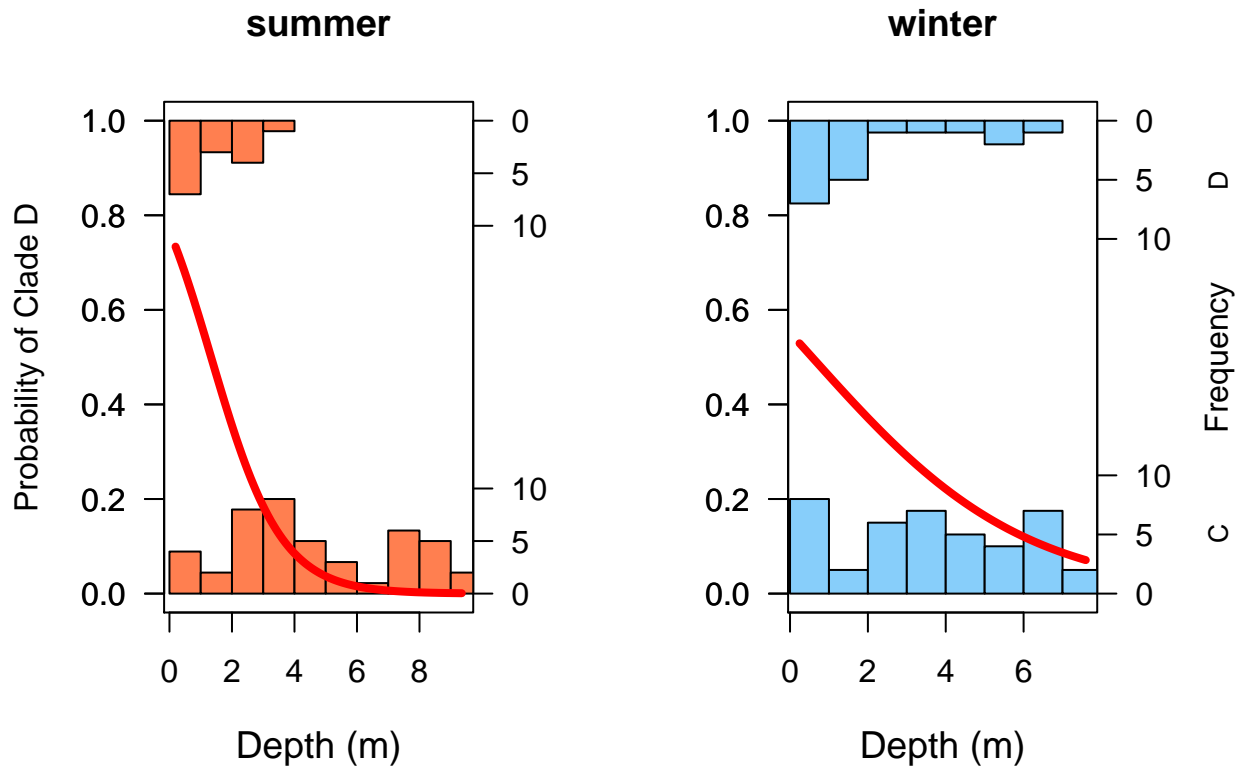
#####
#####
par(mfrow=c(1,2))
Dom1 <- subset(Symb, !is.na(newDepth) & !is.na(Dominant))

```

```
Dom.sum<-Dom1[(Dom1$Season=="summer"),]
Dom.win<-Dom1[(Dom1$Season=="winter"),]

logi.hist.plot(Dom.sum$newDepth, Dom.sum$Dominant, boxp = FALSE, type = "hist", col="coral", xlabel = "Depth (m)",
mtext(side = 2, text = "Probability of Clade D", line = 3, cex = 1)

logi.hist.plot(Dom.win$newDepth, Dom.win$Dominant, boxp = FALSE, type = "hist", col="lightskyblue", xlabel = "Depth (m)",
mtext(side = 4, text = "Frequency", line = 0.5, cex=1)
mtext(side = 4, text = "C", line = 0.5, cex = 0.8)
mtext(side = 4, text = "D", line = 0.5, cex = 0.8)
```



```
dev.copy(pdf, "figures/symbionts/Symbionts_Season_logistic.pdf", encod="MacRoman", height=5, width=8)
dev.off()
```

## Physiology

```
#####
# Physiology
df.fig<-data.trim
# figure layout
layout(matrix(c(1,1,2,3), 2,2, byrow=TRUE))
par(mar=c(5,4.5,2,2))

#####
#####

## season relationship
data.summer<-df.fig[df.fig$Season=="summer", ]
data.winter<-df.fig[df.fig$Season=="winter", ]
```

```

## Site relationship
F8.R10.df<-df.fig[df.fig$Location=="F8-R10", ]
F1.R46.df<-df.fig[df.fig$Location=="F1-R46", ]
HIMB.df<-df.fig[df.fig$Location=="HIMB", ]
R42.df<-df.fig[df.fig$Location=="R42", ]

## Symbiont relationship
C.sum.df<-df.fig[(df.fig$Dom=="C" & df.fig$Season=="summer") ,]
C.win.df<-df.fig[(df.fig$Dom=="C" & df.fig$Season=="winter") ,]; C.win.df<-na.omit(C.win.df)
D.sum.df<-df.fig[(df.fig$Dom=="D" & df.fig$Season=="summer") ,]; D.sum.df<-na.omit(D.sum.df)
D.win.df<-df.fig[(df.fig$Dom=="D" & df.fig$Season=="winter") ,]; D.win.df<-na.omit(D.win.df)

#####
# Fig: chlorophyll (total) over season and depth
#####

plot(chltot~newDepth, data=data.winter,col="dodgerblue3", pch=16, cex=0.7,
     xlab="Depth (m)", ylab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")),
     main="chl: Depth and Season",
     xlim=c(0,11), ylim=c(1,17))
abline((lm(chltot~newDepth, data=data.winter)), col='dodgerblue3', lwd=2)
points(chltot~newDepth, data=data.summer, col="tomato2", pch=16, cex=0.7)
abline((lm(chltot~newDepth, data=data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
     col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.08, 0.1))

##### density plot: by season
plot(density(data.winter$chltot), ylim=c(0,0.3), xlim=c(0, 18), col="dodgerblue3", main="chl: Seasons",
     xlab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")))
lines(density(data.summer$chltot), col="tomato2")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
     col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.35, 0.1))

##### density plot: by Site
plot(density(F1.R46.df$chltot), ylim=c(0,0.3), xlim=c(0, 18), col="tomato2", main="chl: Sites",
     xlab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")), ylab="Density")
lines(density(R42.df$chltot), col="skyblue3")
lines(density(F8.R10.df$chltot), col="springgreen4")
lines(density(HIMB.df$chltot), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
     col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))
dev.copy(pdf, "figures/physiology/PanKB.chl.pdf", width=9, height=7)

#####
# Fig: chlorophyll (total) over season and depth
#####

plot(chltot~Light, data=C.sum.df,col="tomato2", pch=16, cex=0.7,
     xlab="Light (DLI)", ylab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")),
     main="chl: Light and Season",

```



```

    xlim=c(0,20), ylim=c(1,12))
abline((lm(chltot~Light, data=C.sum.df)), col='tomato2', lwd=2)
points(chltot~Light, data=C.win.df, col='dodgerblue3', pch=16, cex=0.7)
abline((lm(chltot~Light, data=C.win.df)), col='dodgerblue3', lwd=2)
points(chltot~Light, data=D.sum.df, col='mediumseagreen', pch=16, cex=0.7)
abline((lm(chltot~Light, data=D.sum.df)), col='mediumseagreen', lwd=2)
points(chltot~Light, data=D.win.df, col='orchid', pch=16, cex=0.7)
abline((lm(chltot~Light, data=D.win.df)), col='orange', lwd=2)
legend("topright", c("C-sum", "C-win", "D-sum", "D-win"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3", "mediumseagreen", "orange"), cex=1, y.intersp = 0.3, x.intersp = 0.3)

plot(density(C.sum.df$chltot), col="tomato2", pch=16, cex=0.7,
     xlab="Light (DLI)", ylab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")),
     main="chl: Light and Season", ylim=c(0, 0.4), xlim=c(0,15), lwd=2)
lines(density(C.win.df$chltot), col="dodgerblue3", lwd=2)
lines(density(D.sum.df$chltot), col="orange", lwd=2)
lines(density(D.win.df$chltot), col="mediumseagreen", lwd=2)
legend("topright", c("C-sum", "C-win", "D-sum", "D-win"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3", "orange", "mediumseagreen"), cex=1, y.intersp = 0.3, x.intersp = 0.3)

##### density plot: by season
plot(density(data.winter$chltot), ylim=c(0,0.3), xlim=c(0, 18), col="dodgerblue3", main="chl: Seasons",
     xlab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")))
lines(density(data.summer$chltot), col="tomato2")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.35, 0.1))

##### density plot: by Site
plot(density(F1.R46.df$chltot), ylim=c(0,0.3), xlim=c(0, 18), col="tomato2", main="chl: Sites",
     xlab=expression(paste("chlorophyll", ~(\mu*g~cm^-2), sep="")), ylab="Density")
lines(density(R42.df$chltot), col="skyblue3")
lines(density(F8.R10.df$chltot), col="springgreen4")
lines(density(HIMB.df$chltot), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))
dev.copy(pdf, "figures/physiology/PanKB.chl.pdf", width=9, height=7)

#####
# Fig: chlorophyll/cell over season and depth
#####
plot(chlcell~newDepth, data=data.winter, col="dodgerblue3", pch=16, cex=0.7,
     xlab="Depth (m)",
     ylab=expression(paste("pg chlorophyll cell"^-1, sep="")), main="chl/cell: Depth and Season",
     xlim=c(0,10), ylim=c(0,15))
abline((lm(chlcell~newDepth, data=data.winter)), col='dodgerblue3', lwd=2)
points(chlcell~newDepth, data=data.summer, col="tomato2", pch=16, cex=0.7)
abline((lm(chlcell~newDepth, data=data.summer)), col='tomato2', lwd=2)

```

```

legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.08, -0.35))

##### density plot: by season
plot(density(data.winter$chlcell), ylim=c(0,0.4), xlim=c(0, 15), col="dodgerblue3", main="chl/cell: Season",
      lines(density(data.summer$chlcell), col="tomato2"))
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.35, -0.08))

##### density plot: by Site
plot(density(F1.R46.df$chlcell), ylim=c(0,0.4), xlim=c(0,15), col="tomato2", main="chl/cell: Sites",
      lines(density(R42.df$chlcell), col="skyblue3")
      lines(density(F8.R10.df$chlcell), col="springgreen4")
      lines(density(HIMB.df$chlcell), col="purple"))
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/physiology/PanKB.chlcell.pdf", width=9, height=7)

#####
# Fig: symbionts over season and depth
#####

plot((zoox/106)~newDepth, data=data.winter, col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab=(expression(paste(~italic("Symbiodinium") ~ (106~cells~cm-2), sep="")))),
      main="zoox: Depth and Season")
abline((lm((zoox/106)~newDepth, data=data.winter)), col='dodgerblue3', lwd=2)
points((zoox/106)~newDepth, data=data.summer, col="tomato2", pch=16, cex=0.7)
abline((lm((zoox/106)~newDepth, data=data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, -0.35))

##### density plot: by season
plot(density(data.summer$zoox/106), col="tomato2", main="zoox: Seasons",
      xlab=(expression(paste(~italic("Symbiodinium") ~ (106~cells~cm-2), sep=""))), xlim=c(0, 7))
lines(density(data.winter$zoox/106), col="dodgerblue3")
legend('topright', c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, -0.08))

##### density plot: by Site
plot(density(F1.R46.df$zoox/106), col="tomato2", main="zoox: Sites",
      xlab=(expression(paste(~italic("Symbiodinium") ~ (106~cells~cm-2), sep=""))), xlim=c(0, 7))
lines(density(R42.df$zoox/106), col="skyblue3")
lines(density(F8.R10.df$zoox/106), col="springgreen4")
lines(density(HIMB.df$zoox/106), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/physiology/PanKB.zoox.pdf", width=9, height=7)

```

```
#####
# Fig: biomass over season and depth
#####

plot(biomass~newDepth, data=data.winter,col="dodgerblue3", pch=16, cex=0.7,
     xlab="Depth (m)", ylab = expression(paste("Total biomass", ~(\text{mg}\cdot\text{cm}^{-2}), sep="")),
     xlim=c(0,10), ylim=c(5,60),
     main="biomass: Depth and Season")
abline(lm(biomass~newDepth, data=data.winter)), col='dodgerblue3', lwd=2)
points(biomass~newDepth, data=data.summer, col="tomato2", pch=16, cex=0.7)
abline(lm(biomass~newDepth, data=data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.08, 0.08))

##### density plot: by season
plot(density(data.winter$biomass), ylim=c(0,0.08), xlim=c(0, 70), col="dodgerblue3", main="biomass: Season")
lines(density(data.summer$biomass), col="tomato2")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), cex=1, y.intersp = 0.3, x.intersp = 0.4, bty="n", inset=c(-0.35, 0.08))

##### density plot: by Site
plot(density(F1.R46.df$biomass), ylim=c(0,0.1), xlim=c(0,70), col="tomato2", main="biomass: Sites",
     xlab=expression(paste("Total biomass", ~(\text{mg}\cdot\text{cm}^{-2}), sep="")))
lines(density(R42.df$biomass), col="skyblue3")
lines(density(F8.R10.df$biomass), col="springgreen4")
lines(density(HIMB.df$biomass), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      y.intersp = 0.3, x.intersp = 0.4,
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/physiology/PanKB.biomass.pdf", width=9, height=7)
```

## Isotopes

```
### ISOTOPES ##
#####

#####
# Fig: d13C host over season and depth
#####

plot(host..d13C~newDepth, data=na.omit(data.winter),col="dodgerblue3", pch=16, cex=0.7,
     xlab="Depth (m)", ylab = expression(paste(delta^{13}, \text{C}[\text{H}], " (\u2030, V-PDB)")),
     xlim=c(0,10), ylim=c(-20,-10),
     main="d13C host: Depth and Season")
abline(lm(host..d13C~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(host..d13C~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(host..d13C~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, 0.08))
```

```
##### density plot: by season
plot(density(na.omit(data.summer$host..d13C)), col="tomato2", main="d13C-host: Seasons",
      xlab=expression(paste(delta~{13}, C[H], " (\u2030, V-PDB))), xlim=c(-20, -8))
lines(density(na.omit(data.winter$host..d13C)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, 0.1))

##### density plot: by Site
plot(density(na.omit(F1.R46.df$host..d13C)), col="tomato2", main="d13C-host: Sites",
      xlab=expression(paste(delta~{13}, C[H], " (\u2030, V-PDB))), ylim=c(0,0.4), xlim=c(-20, -8))
lines(density(na.omit(R42.df$host..d13C)), col="skyblue3")
lines(density(na.omit(F8.R10.df$host..d13C)), col="springgreen4")
lines(density(na.omit(HIMB.df$host..d13C)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d13C-host.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: d13C symb over season and depth
#####

plot(symb..d13C~newDepth, data=na.omit(data.winter), col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab = expression(paste(delta~{13}, C[S], " (\u2030, V-PDB))),
      xlim=c(0,10), ylim=c(-20,-10),
      main= "d13C symb: Depth and Season")
abline(lm(symb..d13C~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(symb..d13C~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(symb..d13C~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, 0.1))

##### density plot: by season
plot(density(na.omit(data.summer$symb..d13C)), col="tomato2", main="d13C-symb: Seasons",
      xlab=expression(paste(delta~{13}, C[S], " (\u2030, V-PDB))), xlim=c(-20, -8))
lines(density(na.omit(data.winter$symb..d13C)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, 0.1))

##### density plot: by Site
plot(density(na.omit(F1.R46.df$symb..d13C)), col="tomato2", main="d13C-symb: Sites",
      xlab=expression(paste(delta~{13}, C[S], " (\u2030, V-PDB))), ylim=c(0,0.4), xlim=c(-20, -8))
lines(density(na.omit(R42.df$symb..d13C)), col="skyblue3")
lines(density(na.omit(F8.R10.df$symb..d13C)), col="springgreen4")
lines(density(na.omit(HIMB.df$symb..d13C)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d13C-symb.pdf", width=9, height=7, encod="MacRoman")
```

```
#####
# Fig: d13C skeleton over season and depth
#####

plot(d13C..skel~newDepth, data=na.omit(data.winter), col="dodgerblue3", pch=16, cex=0.7,
     xlab="Depth (m)", ylab = expression(paste(delta~{13}, C[Skel], " (\u2030, V-PDB)")),
     xlim=c(0,10), ylim=c(-6,2),
     main= "d13C skel: Depth and Season")
abline(lm(d13C..skel~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(d13C..skel~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(d13C..skel~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, -0.08))

##### density plot: by season
plot(density(na.omit(data.summer$d13C..skel)), col="tomato2", main="d13C-skel: Seasons",
     xlab=expression(paste(delta~{13}, C[Skel], " (\u2030, V-PDB)")), ylim=c(0, 0.5), xlim=c(-8, 6))
lines(density(na.omit(data.winter$d13C..skel)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, -0.35))

##### density plot: by Site
plot(density(na.omit(F1.R46.df$d13C..skel)), col="tomato2", main="d13C-skel: Sites",
     xlab=expression(paste(delta~{13}, C[Skel], " (\u2030, V-PDB)")), ylim=c(0,0.5), xlim=c(-8, 6))
lines(density(na.omit(R42.df$d13C..skel)), col="skyblue3")
lines(density(na.omit(F8.R10.df$d13C..skel)), col="springgreen4")
lines(density(na.omit(HIMB.df$d13C..skel)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d13C-skel.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: d13C host-symb over season and depth
#####

plot(d13C..host.sym~newDepth, data=na.omit(data.winter), col="dodgerblue3", pch=16, cex=0.7,
     xlab="Depth (m)", ylab = expression(paste(delta~{13}, C[H-S], " (\u2030, V-PDB)")),
     xlim=c(0,10), ylim=c(-2.5,3),
     main= "d13C h-s: Depth and Season")
abline(lm(d13C..host.sym~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(d13C..host.sym~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(d13C..host.sym~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, -0.08))

##### density plot: by season
plot(density(na.omit(data.summer$d13C..host.sym)), col="tomato2", main="d13C h-s: Seasons",
     xlab=expression(paste(delta~{13}, C[H-S], " (\u2030, V-PDB)")), ylim=c(0,1.5), xlim=c(-3, 5))
lines(density(na.omit(data.winter$d13C..host.sym)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, -0.35))
```



```
##### density plot: by Site
plot(density(na.omit(F1.R46.df$d13C..host.sym)), col="tomato2", main="d13C h-s: Sites",
      expression(paste(delta~{13}, C[H-S], " (\u2030, V-PDB)")), ylim=c(0,1.7), xlim=c(-3,5))
lines(density(na.omit(R42.df$d13C..host.sym)), col="skyblue3")
lines(density(na.omit(F8.R10.df$d13C..host.sym)), col="springgreen4")
lines(density(na.omit(HIMB.df$d13C..host.sym)), col="purple")
legend("topright",c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d13Ch-s.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: d15N host over season and depth
#####

plot(host..d15N~newDepth, data=na.omit(data.winter),col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab = expression(paste(delta~{15}, N[H], " (\u2030, air)")), xlim=c(0,10), ylim=
      main= "d15N host: Depth and Season")
abline(lm(host..d15N~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(host..d15N~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(host..d15N~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08

##### density plot: by season
plot(density(na.omit(data.summer$host..d15N)), col="tomato2", main="d15N-host: Seasons",
      xlab=expression(paste(delta~{15}, N[H], " (\u2030, air)")), xlim=c(0, 12), ylim=c(0, 1))
lines(density(na.omit(data.winter$host..d15N)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35

##### density plot: by Site
plot(density(na.omit(F1.R46.df$host..d15N)), col="tomato2", main="d15N-host: Sites", xlab=expression(pa
lines(density(na.omit(R42.df$host..d15N)), col="skyblue3")
lines(density(na.omit(F8.R10.df$host..d15N)), col="springgreen4")
lines(density(na.omit(HIMB.df$host..d15N)), col="purple")
legend("topright" ,c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d15N-host.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: d15N symb over season and depth
#####

plot(symb..d15N~newDepth, data=na.omit(data.winter),col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab =expression(paste(delta~{15}, N[S], " (\u2030, air)")),
      xlim=c(0,10), ylim=c(2,7),
      main= "d15N symb: Depth and Season")
```

```

abline(lm(symb..d15N~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(symb..d15N~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(symb..d15N~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, 0.08))

##### density plot: by season
plot(density(na.omit(data.summer$symb..d15N)), col="tomato2", main="d15N-symb: Seasons",
      xlab= expression(paste(delta~{15}, N[S], " (\u2030, air)")), xlim=c(0, 9), ylim=c(0, 0.9))
lines(density(na.omit(data.winter$symb..d15N)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, 0.08))

##### density plot: by Site
plot(density(na.omit(F1.R46.df$symb..d15N)), col="tomato2", main="d15N-symb: Sites",
      xlab= expression(paste(delta~{15}, N[S], " (\u2030, air)")), xlim=c(0, 9), ylim=c(0,1.8))
lines(density(na.omit(R42.df$symb..d15N)), col="skyblue3")
lines(density(na.omit(F8.R10.df$symb..d15N)), col="springgreen4")
lines(density(na.omit(HIMB.df$symb..d15N)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d15N-symb.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: d15N host.symb over season and depth
#####

plot(d15N..host.sym~newDepth, data=na.omit(data.winter),col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab = expression(paste(delta~{15}, N[H-S], " (\u2030, air)")),
      xlim=c(0,10), ylim=c(-1,2),
      main= "d15N h-s: Depth and Season")
abline(lm(d15N..host.sym~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(d15N..host.sym~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(d15N..host.sym~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08, 0.08))

##### density plot: by season
plot(density(na.omit(data.summer$d15N..host.sym)), col="tomato2", main="d15N h-s: Seasons",
      xlab=expression(paste(delta~{15}, N[H-S], " (\u2030, air)")), xlim=c(-1, 3), ylim=c(0,2))
lines(density(na.omit(data.winter$d15N..host.sym)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.3, cex=1, bty="n", inset=c(-0.35, 0.08))

##### density plot: by Site
plot(density(na.omit(F1.R46.df$d15N..host.sym)), col="tomato2", main="d15N h-s: Sites", xlab=expression
lines(density(na.omit(R42.df$d15N..host.sym)), col="skyblue3")
lines(density(na.omit(F8.R10.df$d15N..host.sym)), col="springgreen4")
lines(density(na.omit(HIMB.df$d15N..host.sym)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),

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```

        col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.d15Nh-s.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: C.N host season and depth
#####
plot(host..C.N~newDepth, data=na.omit(data.winter),col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab = expression(paste("C:N" [H])),
      xlim=c(0,10), ylim=c(5,10),
      main= "C:N-host Depth and Season")
abline(lm(host..C.N~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(host..C.N~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(host..C.N~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08

##### density plot: by season
plot(density(na.omit(data.summer$host..C.N)), col="tomato2", main="C:N-host Depth and Season",
      xlab=expression(paste("C:N" [H])), ylim=c(0, 1), xlim=c(4, 10))
lines(density(na.omit(data.winter$host..C.N)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35

##### density plot: by Site
plot(density(na.omit(F1.R46.df$host..C.N)), col="tomato2", main="C:N-host Depth and Season",
      xlab=expression(paste("C:N" [H])), ylim=c(0,1.5), xlim=c(4, 10))
lines(density(na.omit(R42.df$host..C.N)), col="skyblue3")
lines(density(na.omit(F8.R10.df$host..C.N)), col="springgreen4")
lines(density(na.omit(HIMB.df$host..C.N)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.C.Nhost.pdf", width=9, height=7, encod="MacRoman")

#####
# Fig: C.N symb season and depth
#####
plot(symb..C.N~newDepth, data=na.omit(data.winter),col="dodgerblue3", pch=16, cex=0.7,
      xlab="Depth (m)", ylab = expression(paste("C:N" [S])),
      xlim=c(0,10), ylim=c(5,14),
      main= "C:N-symb Depth and Season")
abline(lm(symb..C.N~newDepth, data=na.omit(data.winter)), col='dodgerblue3', lwd=2)
points(symb..C.N~newDepth, data=na.omit(data.summer), col="tomato2", pch=16, cex=0.7)
abline(lm(symb..C.N~newDepth, data=na.omit(data.summer)), col='tomato2', lwd=2)
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.08

##### density plot: by season

```



```

plot(density(na.omit(data.summer$symb..C.N)), col="tomato2", main="C:N-symb Depth and Season",
      xlab=expression(paste("C:N"[S])), ylim=c(0, 0.8), xlim=c(5, 15))
lines(density(na.omit(data.winter$symb..C.N)), col="dodgerblue3")
legend("topright", c("summer", "winter"), lty=c(1,1), lwd=c(2,2),
      col=c("tomato2", "dodgerblue3"), y.intersp = 0.3, x.intersp = 0.4, cex=1, bty="n", inset=c(-0.35, 0.1))

##### density plot: by Site
plot(density(na.omit(F1.R46.df$symb..C.N)), col="tomato2", main="C:N-symb Depth and Season",
      xlab=expression(paste("C:N"[S])), ylim=c(0,0.8), xlim=c(5, 15))
lines(density(na.omit(R42.df$symb..C.N)), col="skyblue3")
lines(density(na.omit(F8.R10.df$symb..C.N)), col="springgreen4")
lines(density(na.omit(HIMB.df$symb..C.N)), col="purple")
legend("topright", c("Fringe-Reef 46", "Patch-Reef 42", "Fringe-Reef 10", "HIMB"), lty=c(1,1), lwd=c(2,2),
      col=c("sienna1", "skyblue3", "springgreen4", "purple"), cex=0.8, bty='n', inset=c(-0.5, -0.1))

dev.copy(pdf, "figures/isotope/PanKB.C.Nsym.pdf", width=9, height=7, encod="MacRoman")

```